

1 agggagagggc agtgaccatg aaggctgtgc tgcttgcccct gttgtaggca
51 ggcttggccc tgcagccagg cactgcccttg ctgtgctact ccttgcaaagc
101 ccaggtgagc aactgaggact gcctgcagg tggagaactgc accccagctgg
151 ggggagcattgctggacccgcgcg cgcarcccgcg cagttggccct cctgacccgtc
201 atcagcaaag gctgcagttg aactgcgtgg atggactcac aggactacta
251 cgtggggcaag aagaacatca cgtgctgtga caccgacttg tgcaacgccca
301 gcgggggccccca tgccctgcag ccggctgccg ccatccttgc gctgctccct
351 gcactcggcc tgctgctctggggacccgggc cagctatagg ctctgggggggg
401 ccccgctgca gcccacactg ggtgggtggc cccaggccttgtgccactc
451 ctcacagaac ctggcccagt gggagccctgt cctggttcct gaggcacatc
501 ctaacgcaag ttgaccaatgtatttgca ccccttttcc ctnaaccctg
551 acttcccatg gggcctttcg caggatccug accttggcagat cagttttag
601 tganacanat ccgcntgcag atggcccttc caaccttttg tgttgttttg
651 tccatggcccc agcatttccc acccttaaaccctgtgtttccagg gcacttttcc
701 ccccaggaaag cttccctgc ccaccccatg tatgaatgta gccaggtttg
751 gtccgttggtg tcccccgcac ccagcagggggg acaggccatcg gggaggggcccc
801 agtaattggct ggatgtaagt ggactggatgta gaactggagggg acaagagtttg
851 acgttggtttc tggggttttttcccagggattg gggccttggggggggcccc
901 gggggccagggc ctccatttttg tggggttcccc gaattggcagccgaccagg
951 cgtaggccctttttttccac cgtttggttta agccaaaaaaaaaaaaaaaa

FIGURE 1A

MKA VLL ALL MAGL ALQPGT ALL CYSCKA QVS NEDCLQV
E NCTQLGEQCWTARIRAVGLLTV I SKGCSLNCVDD S
QDYYVVGKKNITCCDTDLCNASGAHALQPAAAILALLP AL
GLLLWGP GQL

FIGURE 1B

ATGAAGACAGTTTTTATCCTGCTGGCCACCTACTTAGCCCTGCATCCAGGTGCTGCT
 1 TACTTCTGTCAAAAAAAATAGGACGACGGTGGATGAATCGGGACGTAGGTCCACGACGA 60

M K T V F F I L L A T Y L A L H P G A A -

CTGAGTGCTATTGACAGCACAGATGAACAAACAGAGACTGTCATGAATGTACAGAAC
 61 GACGTCACGATAAGTACGTGTCGTCTACTTGTGTCATGACAGACTTACATGCTTG 120

L Q C Y S C T A Q M N N R D C L N V Q N -

TGCAGCCTGGACCAGCACAGTTGCTTACATCGCGATCCGGGCCATTGGACTCGTGACA
 121 ACGTCGGACCTGGTCGTCAACGAAATGTAGCGCGTAGGCCCGTAACCTGAGCACTGT 180

C S L D Q H S C F T S R I R A I G L V T -

'GTTATCAGTAAGGGCTGCAGCTCACAGTGTGAGGATGACTCGGAGAACTACTATTGGGC
 181 CAATAGTCATTCCGACGTCGAGTGTACACTCCTACTGAGCCTTGTATGATAAACCCG 240

V I S K G C S S Q C E D D S E N Y Y L G -

AAGAAAGAACATCACGTGCTGCTACTCTGACCTGTGCAATGTCAACGGGCCACACCTG
 241 TTCTCTTGAGTGCACGACGATGAGACTGGACAGTACAGTTGCCCCGGTGTGGGAC 300

K K N I T C C Y S D L C N V N G A H T L -

AAGGCCACCCACCAACCTGGGCTGCTGACCGTGCTCTGCAGCCTGTTGCTGTGGGCTCC
 301 TTGGTGGTGGTGGGGACCCGACGACTGGCACGAGACGTCGGACAACGACACCCCGAGG 360

K P P T T L G L L T V L C S L L L W G S -

ACCCGTCTGAGGCTCTGGGAGAGCCTACCATAGCCCATTGTGAAGGGATGAGCTGCAC
 361 TCGGGAGACATCCGAGACCCCTCTCGGATGGTATGGCTAACACTCCCTACTCGACGTG 420

S R L *

TCCACCCCCACCCCCACACAGG

421 AGGTGGGGTGGGGTGTGTCC 441

FIGURE 2

1 M K I F P P V I D E A N L I G V E R A S S NSCA-2
1 M K A V I L L A L L M A G G D A K O P G T A. MPSCA
1 M K I V I L S L L Q A T Y M A M H P G A A. MPSCA

21 E M C F S C L N Q K S I N * L Y C E K P T I
21 E L C Y S C K A Q V S N * S D C I L Q V E N *
21 L Q C Y S C T A Q M H N N * R D C L N V Q N *

41 C S O Q O N Y C V T V S A S X G I G N L
41 C T Q L G E Q C W T A R I R A V G L L T
41 C S L O Q H S C F T S R I R A I G L V T

61 V T F G H S L S N T C I S P A C P I P E G
61 V - - - - - I S K G C I S L N C V D D S Q
61 V - - - - - I S K G C I S S Q C E D D S E

81 V N V G V A S M G I S C C Q S F L C N * F
76 D Y Y V G K K - N * L I C C C D T D L C I N S A
76 N Y Y L G K K - N P I T C C Y S D E C N * V

101 S I A A D G G I R A S V T D E G A G G W L
95 S P G A H A E O P T A A A I H A L I P A E G
95 N G A E T Z K P P T T L G G E T V D C S

121 S L T P A L L R F G P
115 L L L W G P G O L - -
115 L L L W G S S R L - -

FIGURE 3

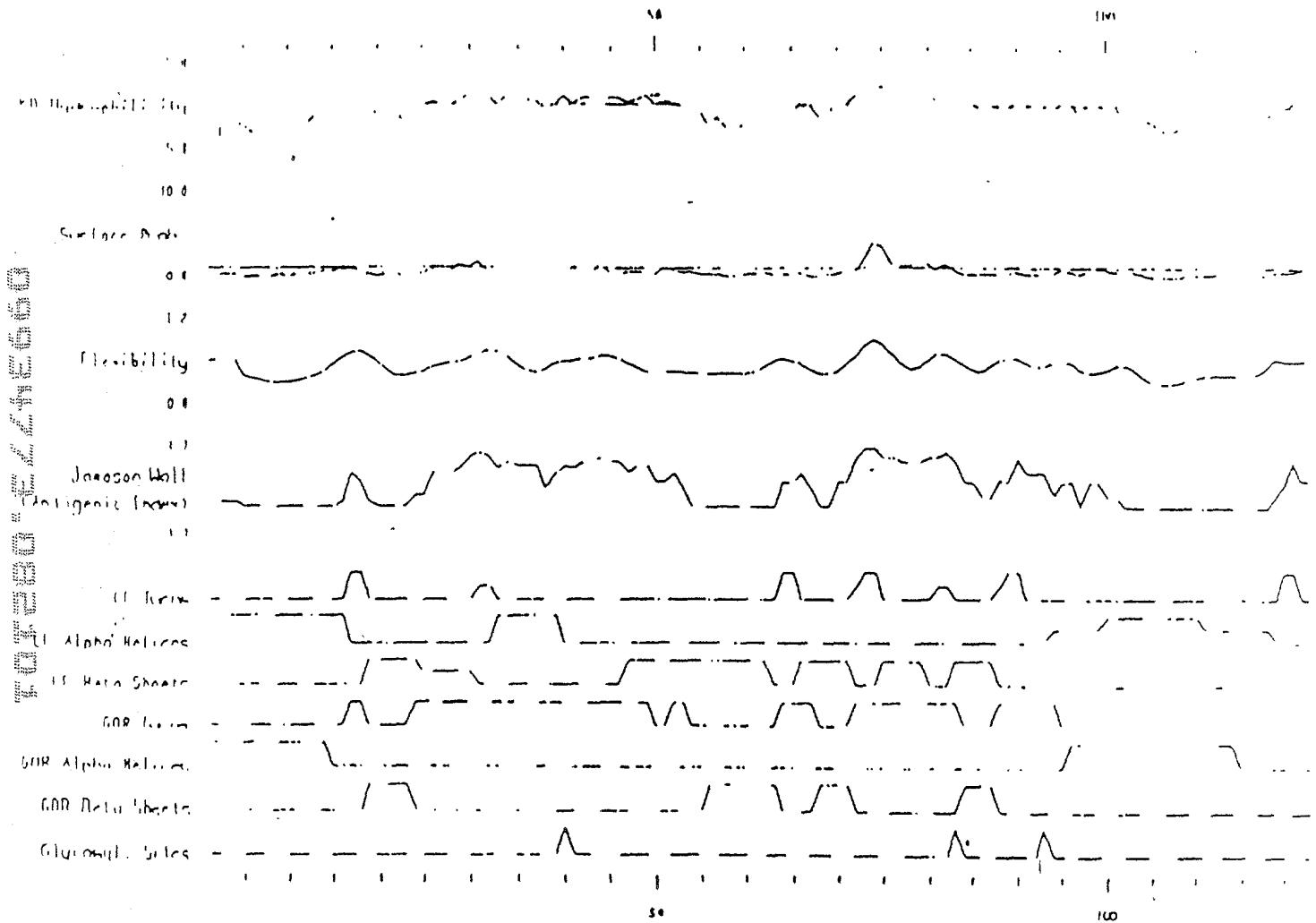


FIGURE 4

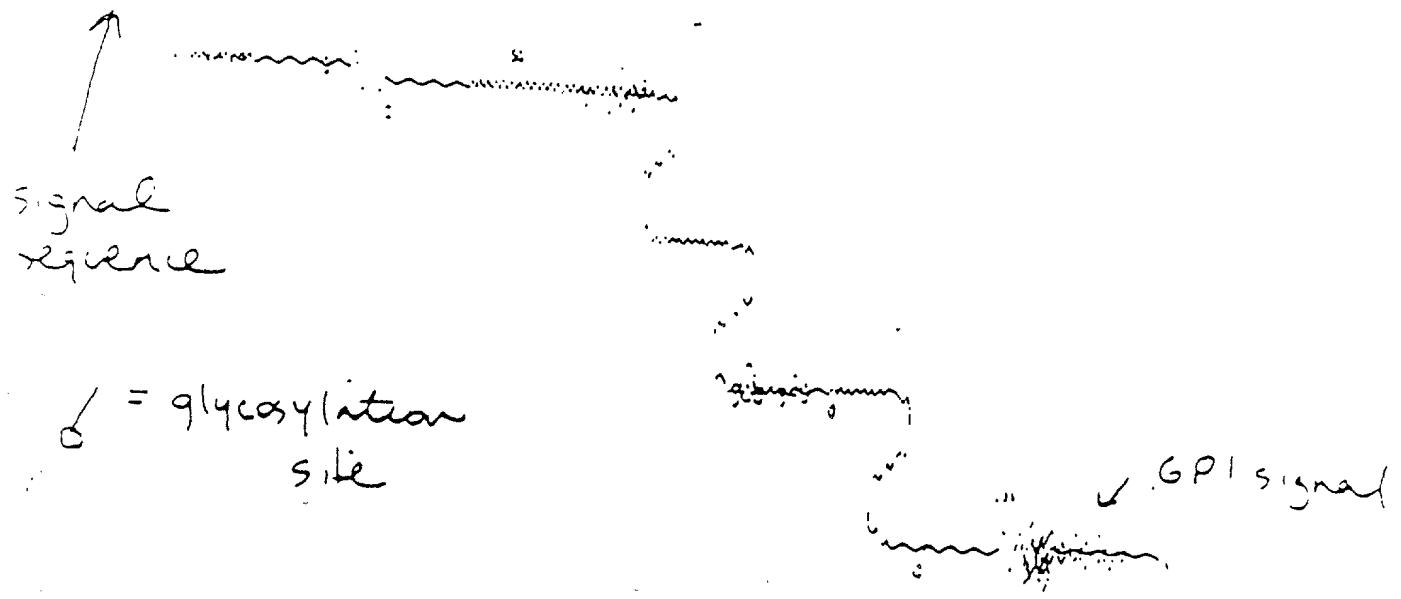
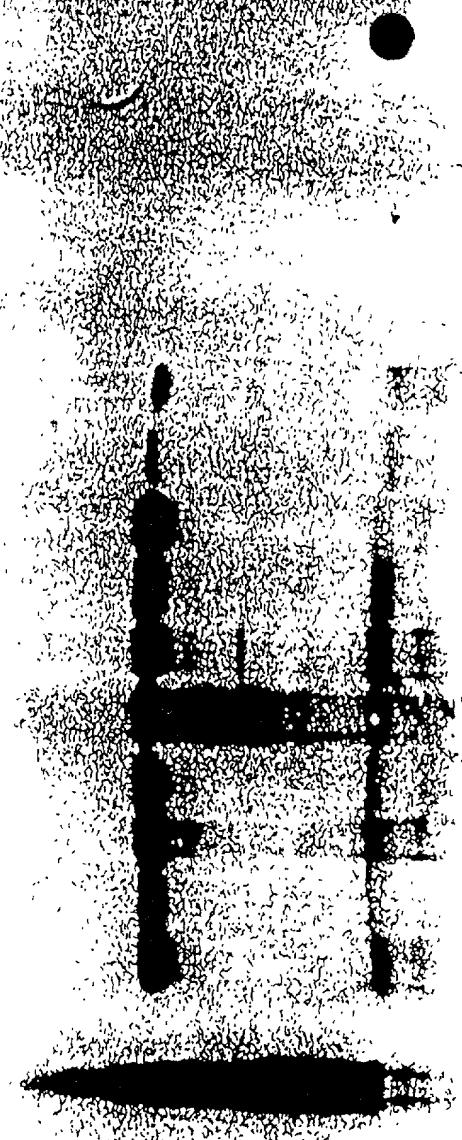


FIGURE 5

Support to be 80%
Normal tissue
Western Blot
1 hr exp

1:1000
168



prostate (Kunz)
prostate (Bukke)
prostate (dck)
Bladder (Kunz)
Bladder (dck)
Bladder (Rob)
Kidney (NL084)
Kidney (NL021)
Fas1
Sm. Intest.

LAPC9

FIGURE 6

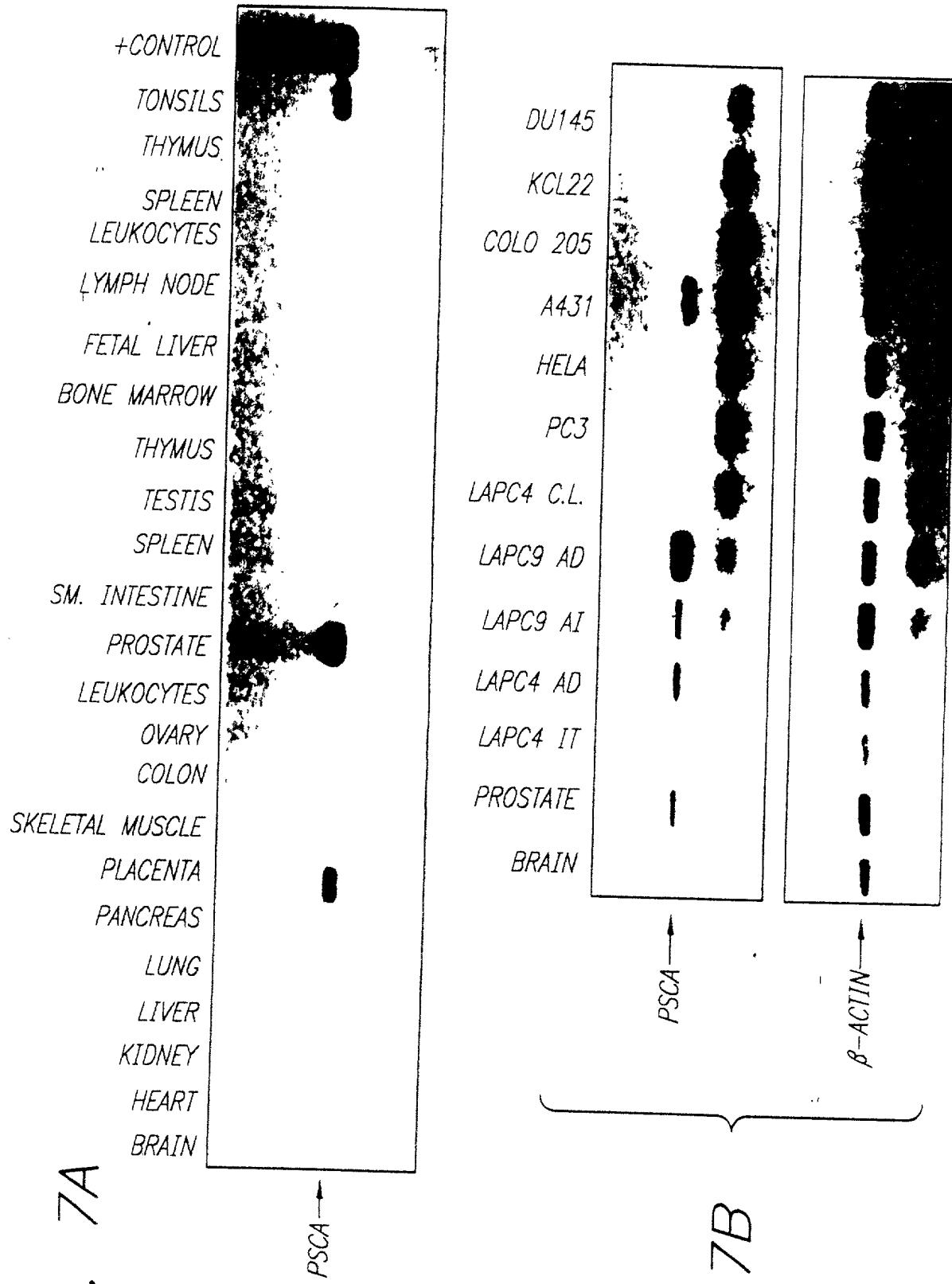


FIG. 7A

FIG. 7B

Legend:  untranslated region of p5CA
 translated region of p5CA

FIG. 8A

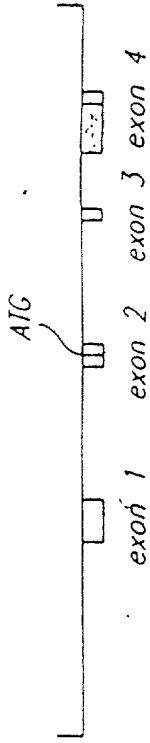


FIG. 8B

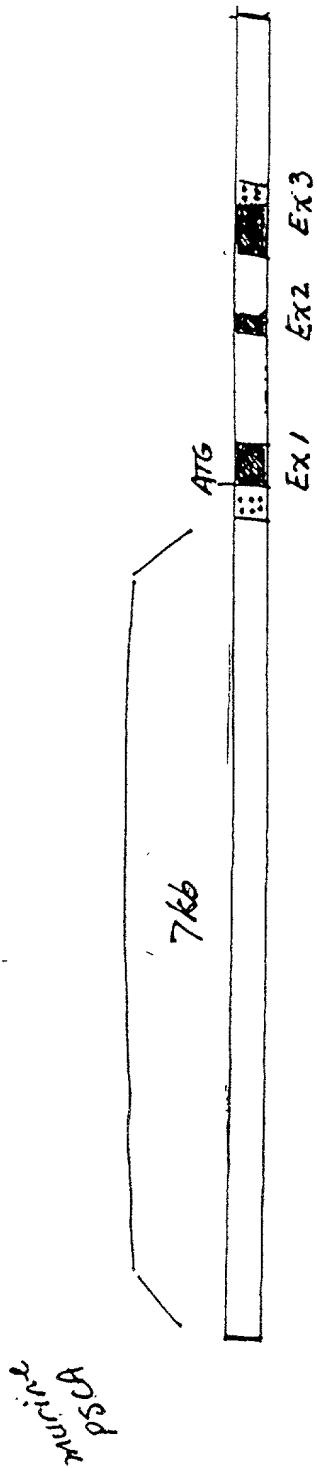


FIG. 8C

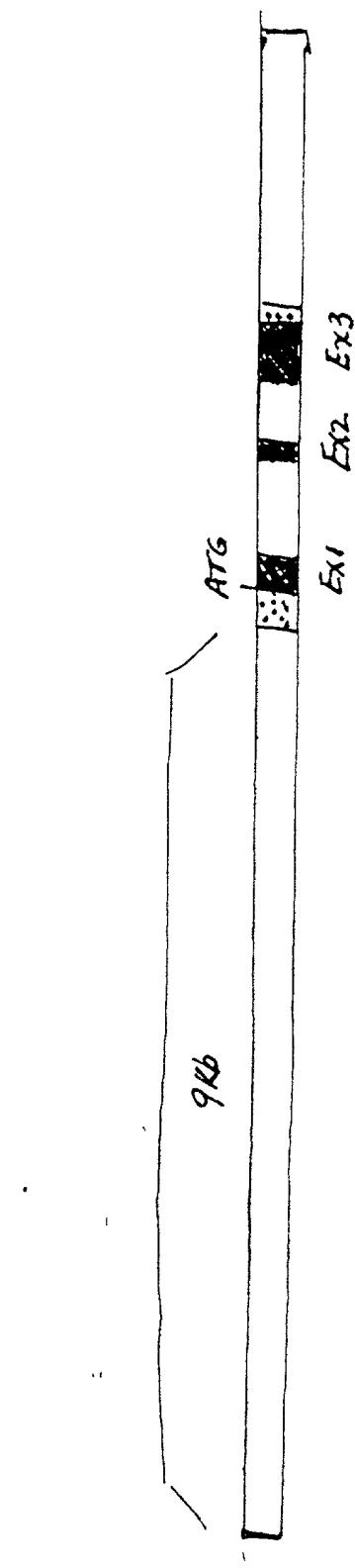


FIGURE 8

murine
p5CA

human
p5CA

100-07207-EE-660

PSCA / PSA Expression in Benign Prostate vs. Prostate Cancer Xenograft

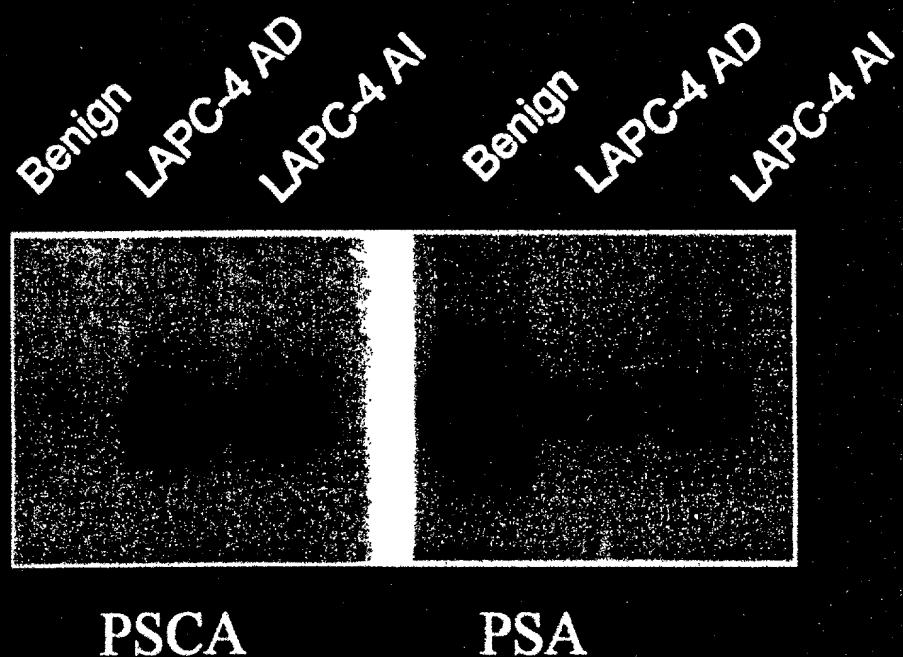
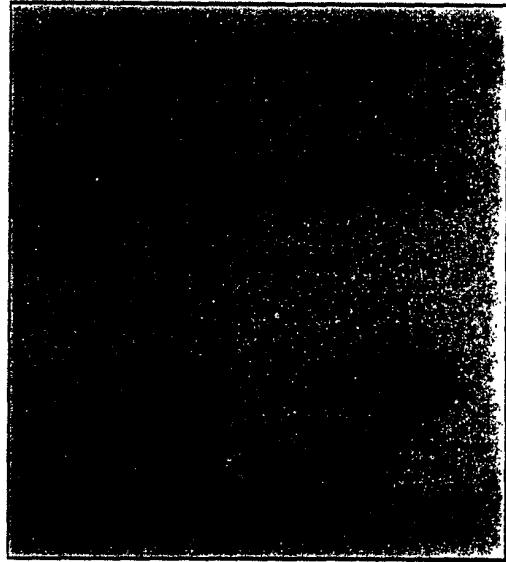


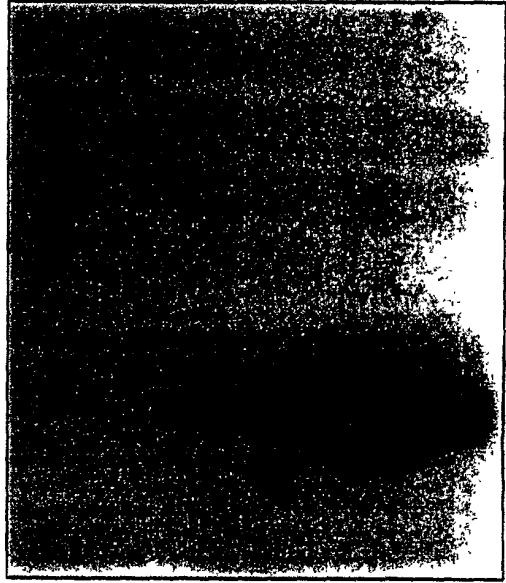
FIGURE 9A

PANCREAS
KIDNEY
SKELETAL MUSCLE
LIVER
LUNG
PLACENTA
BRAIN
HEART



~1kb

PERIPHERAL LEUKOCYTES
COLON
SMALL INTESTINE
OVARY
TESTIS
PROSTATE
THYMUS
SPLEEN



PSCA

FIG. 9B

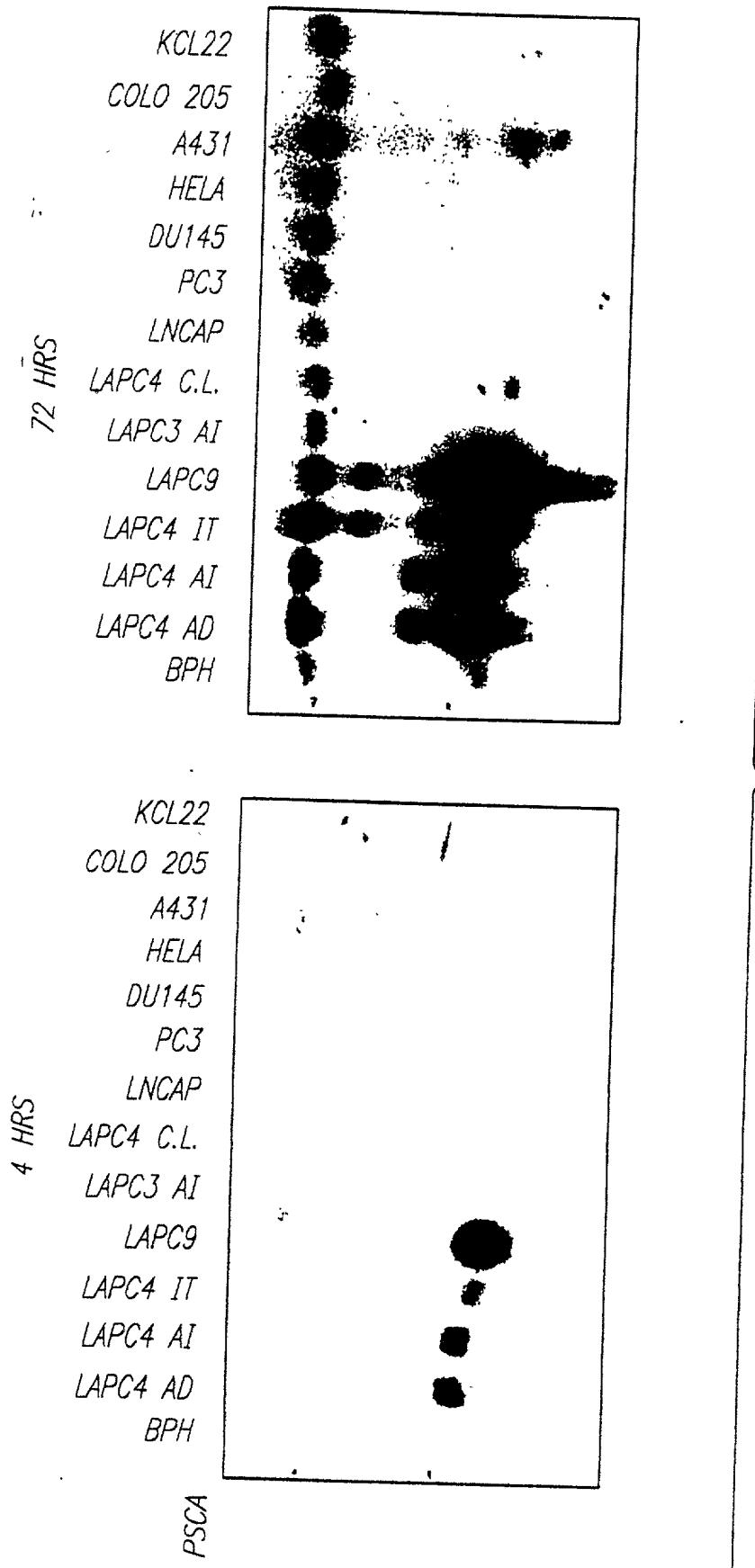


FIG. 10-1

FIG. 10-2

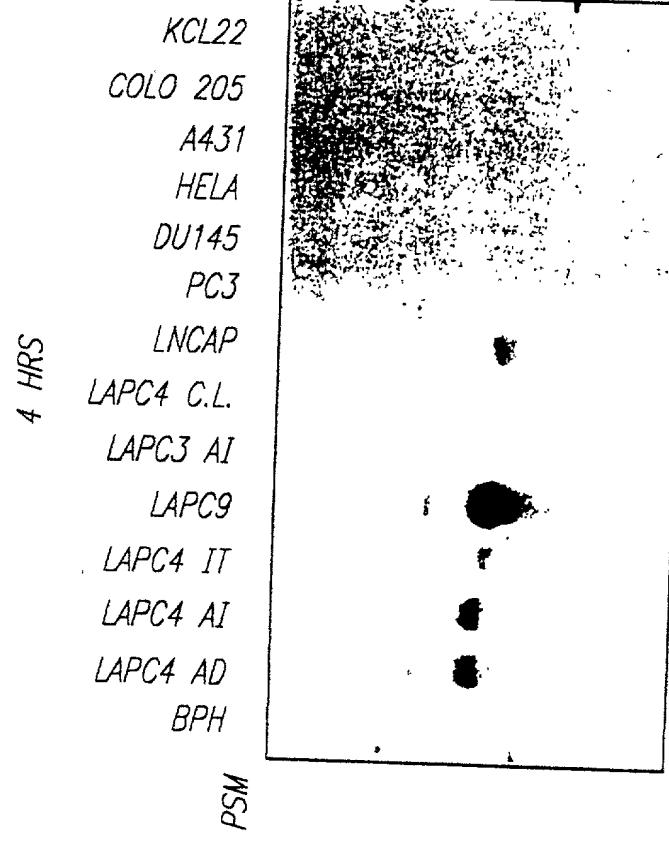
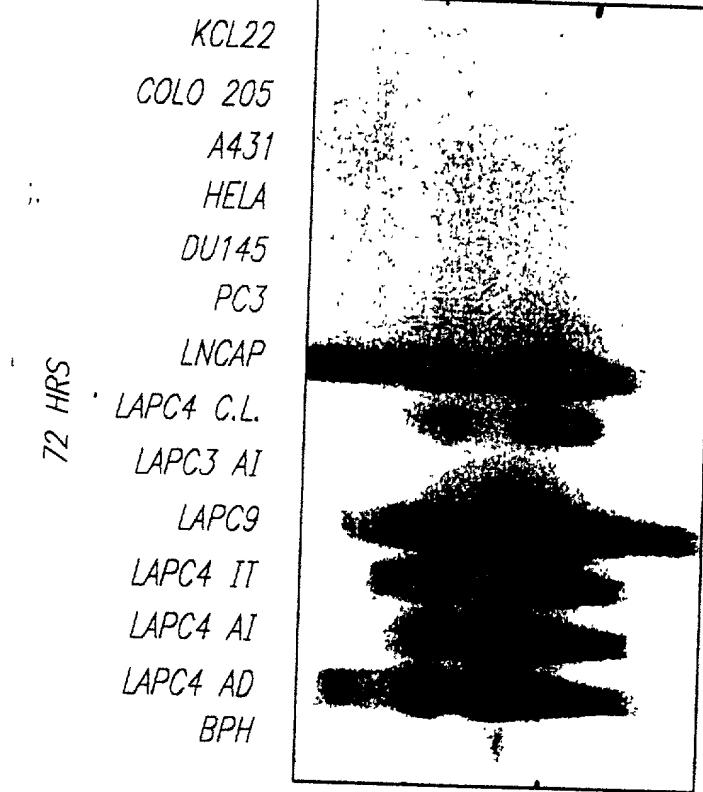
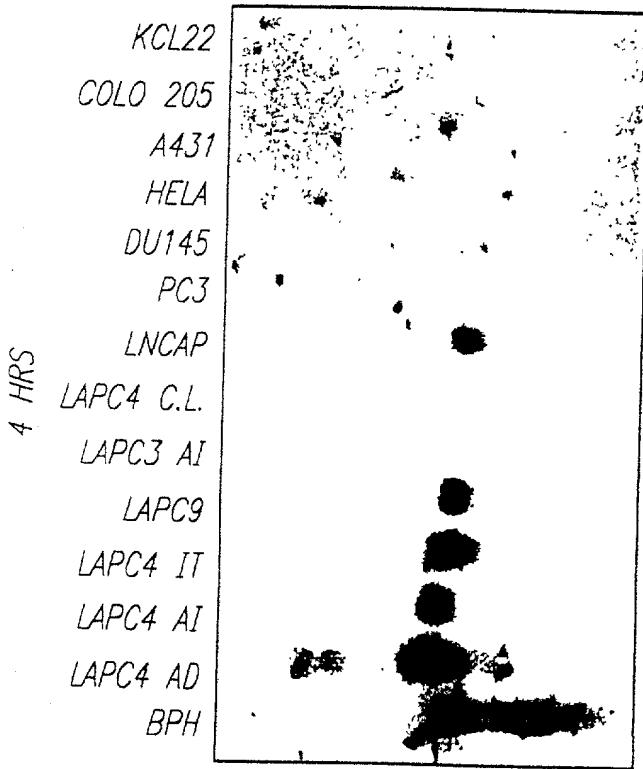
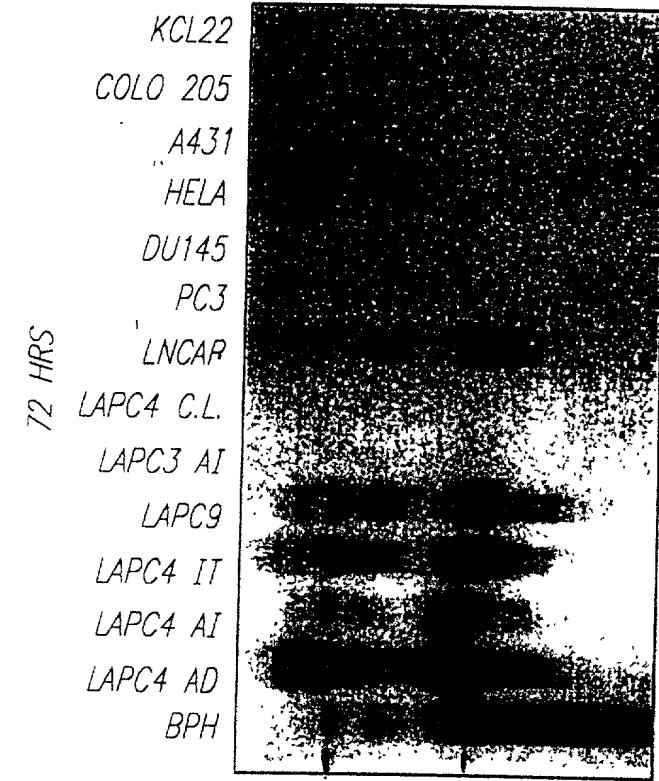


FIG. 10-2



PSA

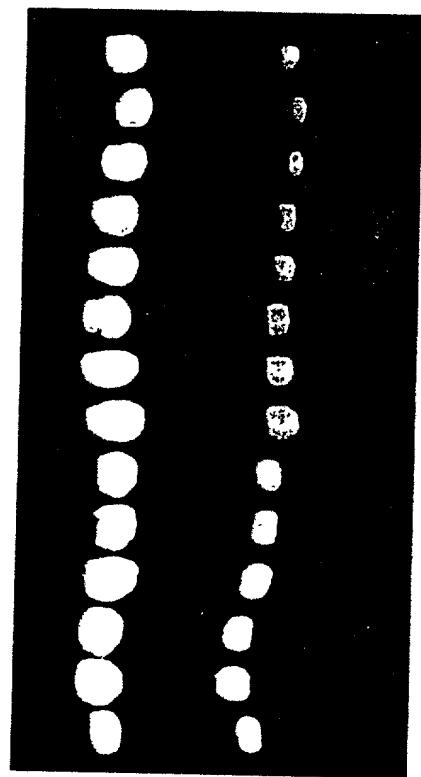


FIG. 10-3

FIG. 11A



புது திட்டங்கள் முன்வரை



FIG. 11B

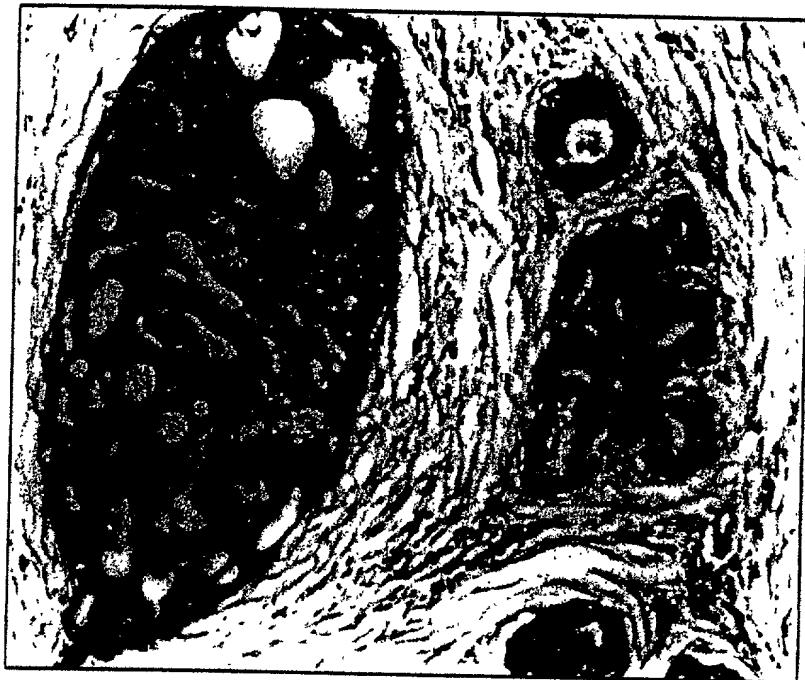


FIG. 11C

FIG. 12A

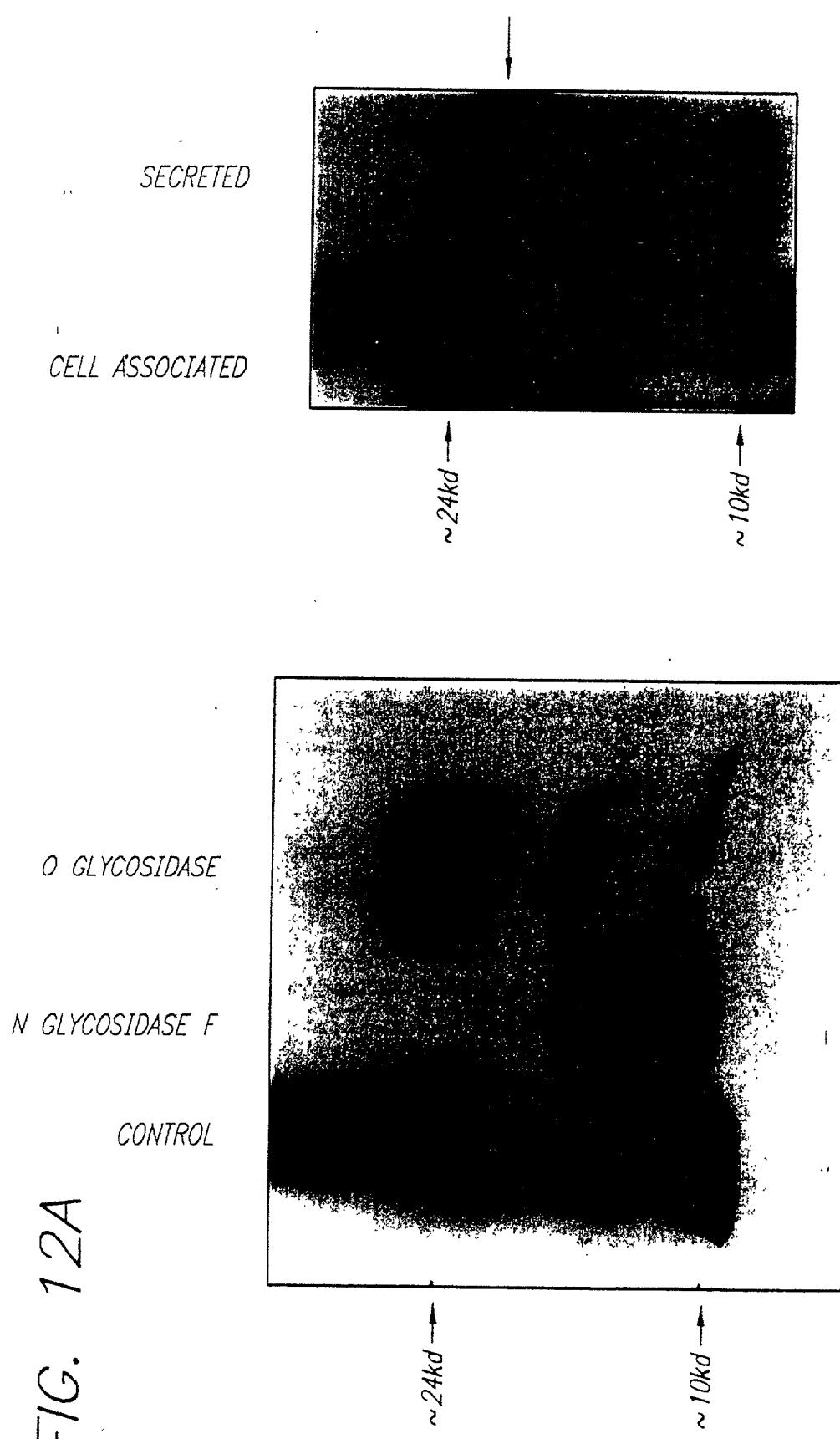


FIG. 12B

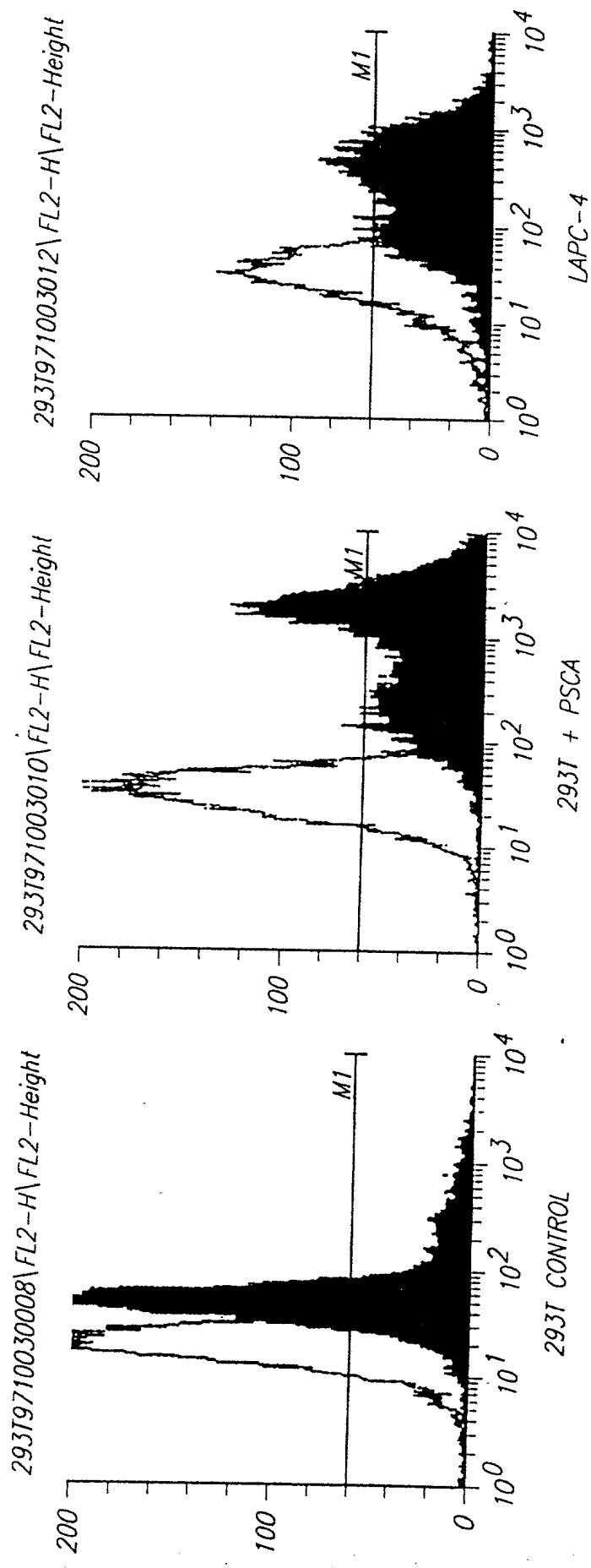
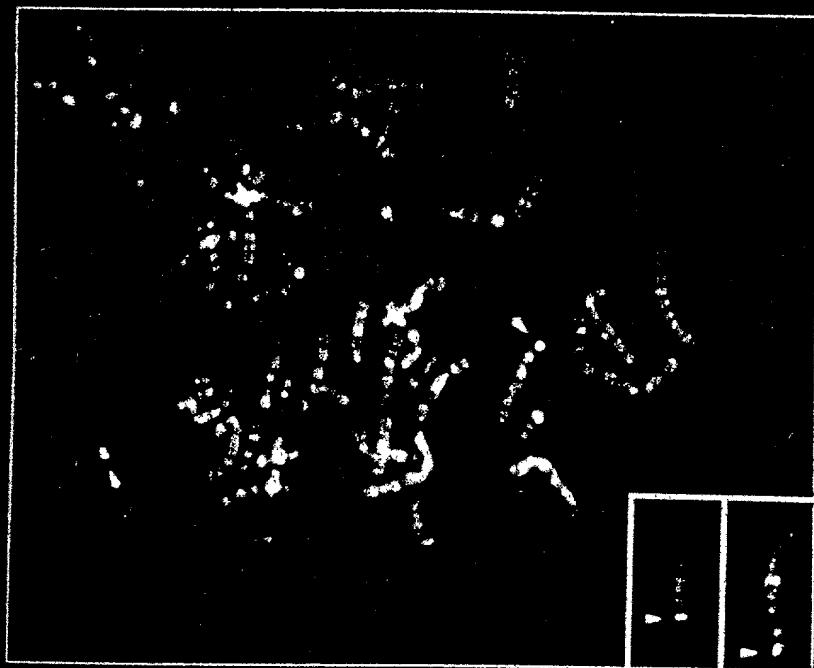


FIGURE 12C

PSCA Maps to Chromosome 8q24.2



Fluorescent
in Situ Hybridization
Analysis of PSCA

FIGURE 13

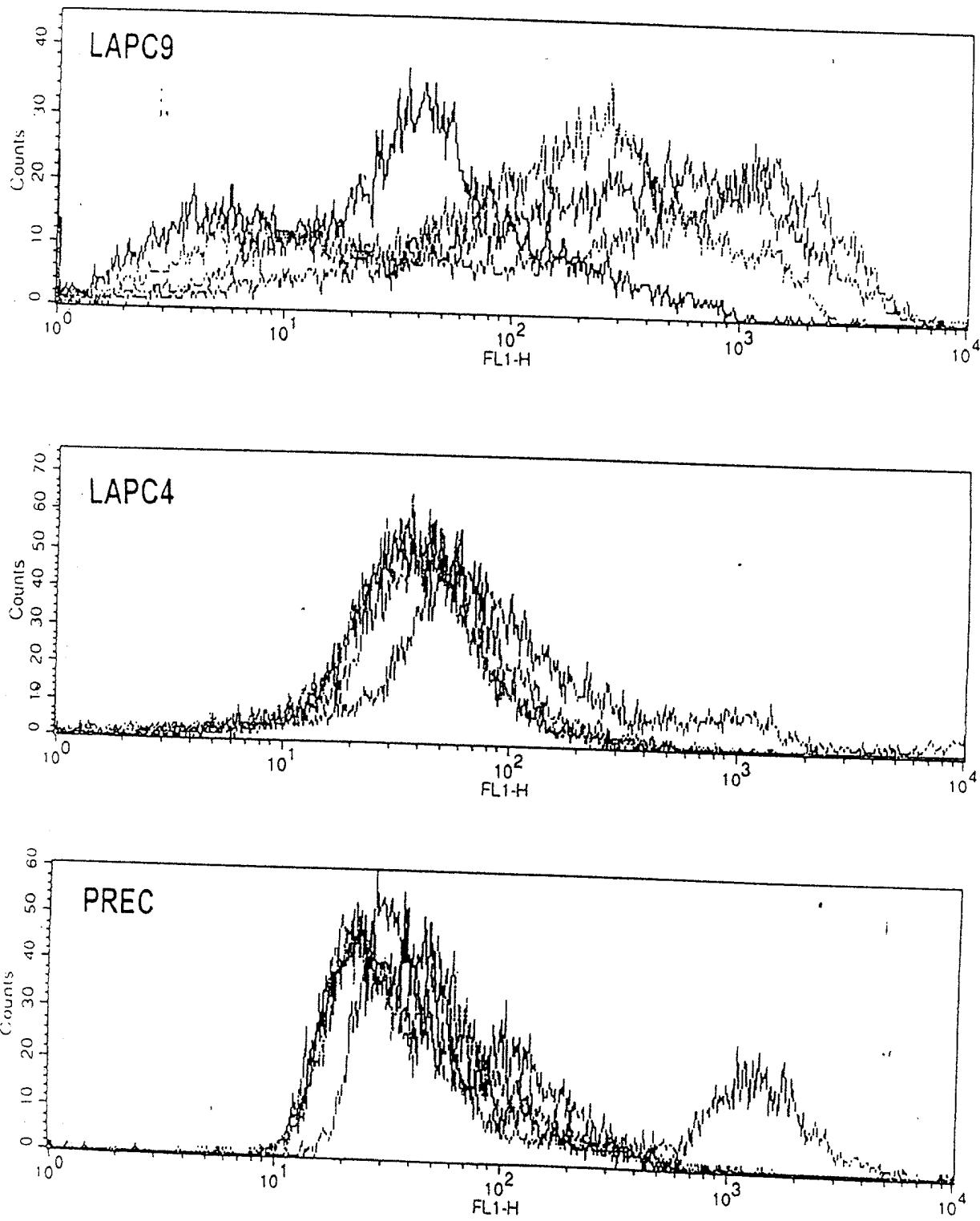


FIGURE 14

A**Epitope map**

mAb	Isotype	EL (18-98)	N (2-50)	M (46-109)	C (85-123)
1G8	IgG1	2.039	0.007	0.628	0.000
2H9	IgG1	1.318	0.863	0.032	0.021
3C5	IgG2a	2.893	1.965	0.016	0.005
3E6	IgG3	0.328	0.024	0.069	0.370
4A10	IgG2a	2.039	1.315	0.000	0.014
2A2	IgG2a	1.366	0.733	0.010	0.003
3G3	IgG2a	2.805	1.731	0.004	0.000

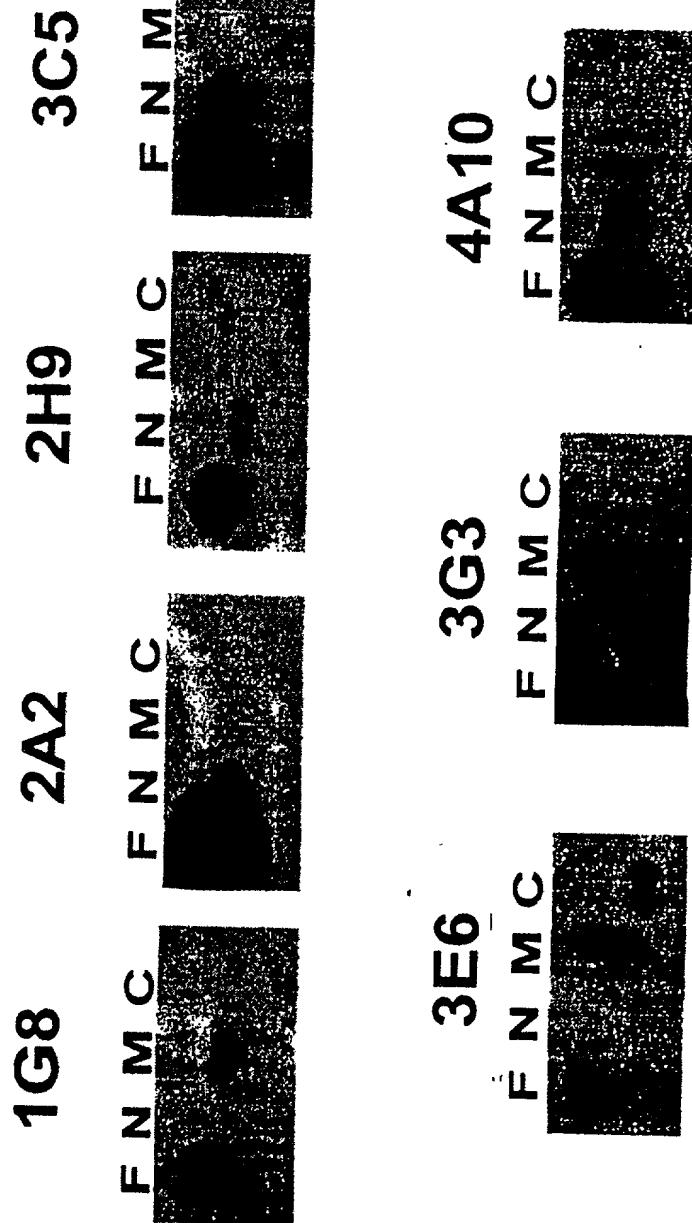
B

FIGURE 15

Prostate Stem Cell Antigen (PSCA) is a GPI-anchored Protein

1	I	F	S	M	P	A	D	E	A	A	L	S	L	A	A	hSCA-2
1	A	A	D	E	A	A	T	E	M	A	G	D	A	L	P	hPSCA
1	T	A	A	D	E	A	T	E	M	A	G	D	A	L	P	mPSCA
21	M	G	C	F	S	C	D	N	Q	S	N*	C	L			
21	G	G	C	F	S	C	D	N	Q	S	N*	D	C	L	V	N*
21	G	G	C	F	S	C	D	N	Q	S	N*	P	D	C	L	N
41	C	F	S	C	D	N	T	A	G	A	A					
41	C	F	S	C	D	N	T	A	G	A	A	R	I	R	A	
41	C	F	S	C	D	N	T	A	G	A	A	R	I	R	A	T
61	V															
61	V															
61	V															
81	V															
81	V															
81	V															
95	S	A	A	D	G	S	-	R	A	A		L	L			
95	S	A	A	D	G	S	-	R	A	A		L	L			
95	S	A	A	D	G	S	-	R	A	A		L	L			
101	S	A	A	D	G	S	-	R	A	A		L	L			
121	S	A	A	D	G	S	-	R	A	A		L	L			
115	S	A	A	D	G	S	-	R	A	A		L	L			
115	S	A	A	D	G	S	-	R	A	A		L	L			

(Reiter, R.E., et al., 1997, *PNAS*)

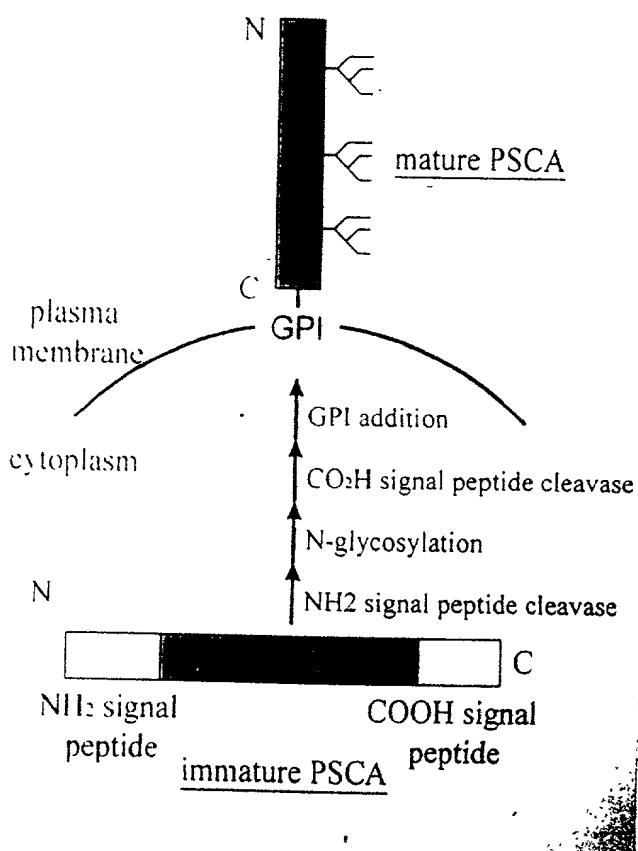
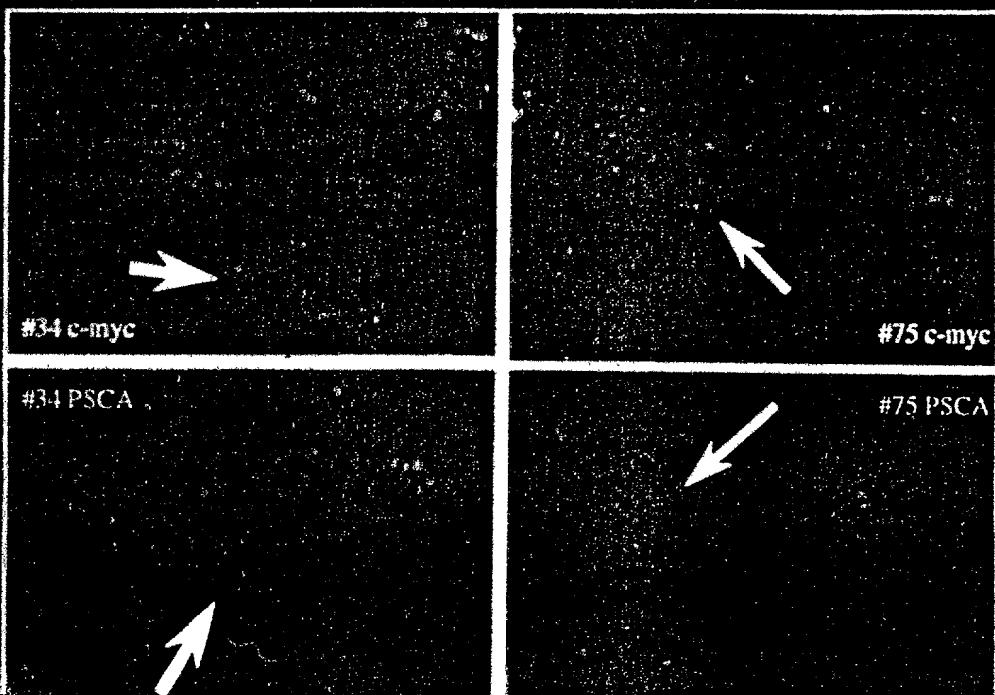


FIGURE 16

FISH Analysis of PSCA and c-myc in Prostate Cancer

Gain Chromosome 8

Amplification



R. Jenkins

FIGURE 17

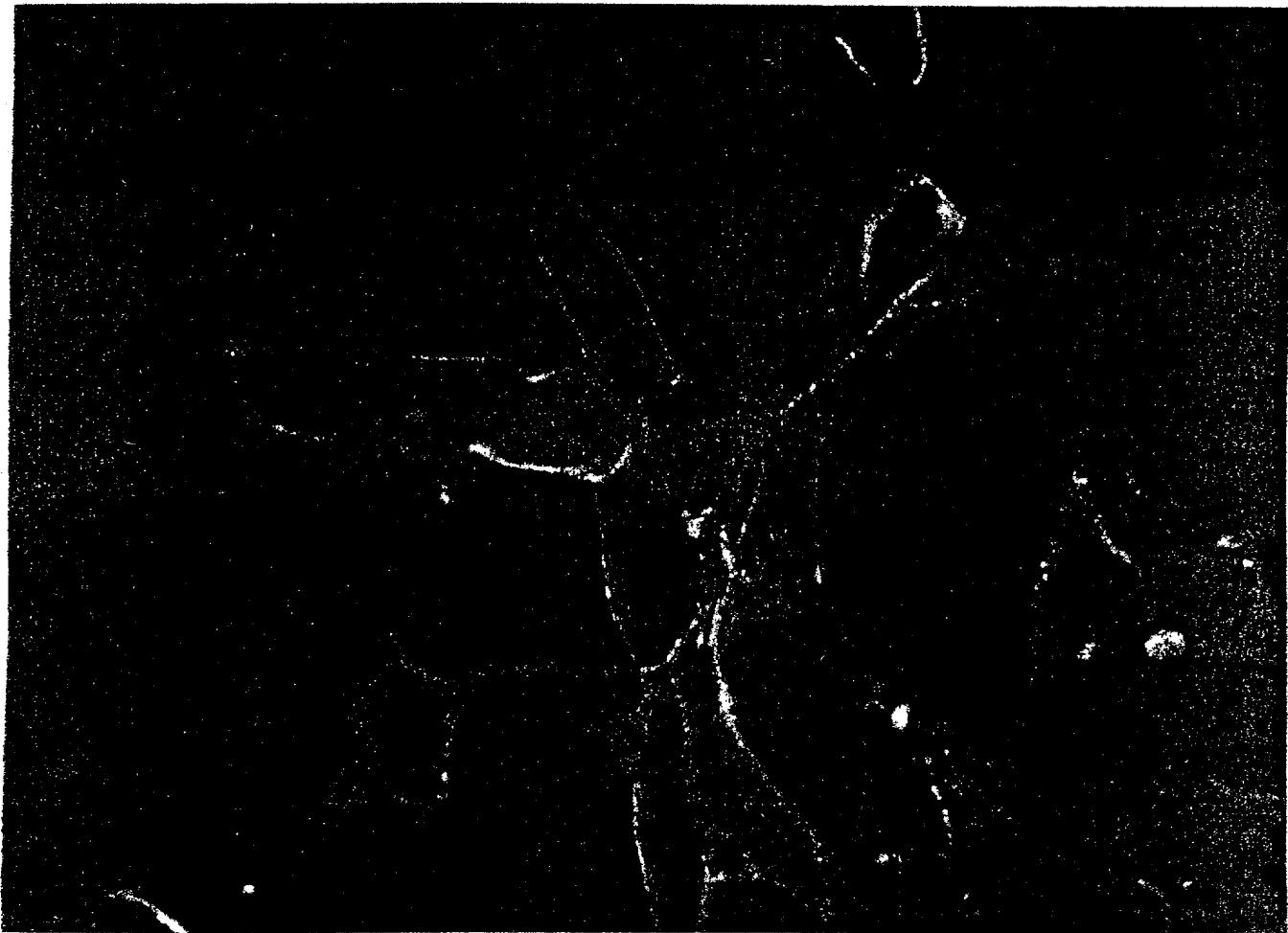


FIGURE 18

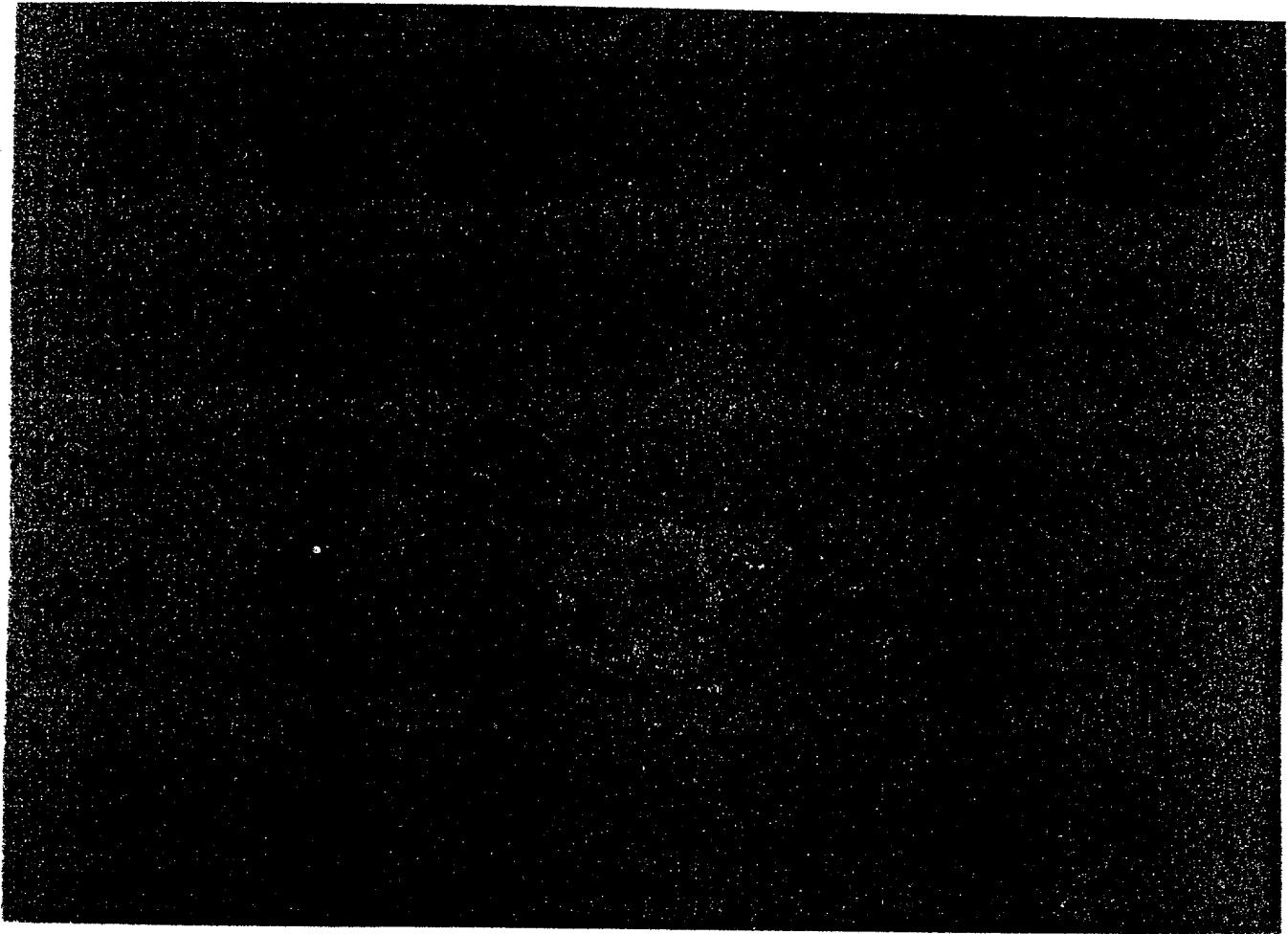


FIGURE 19



FIGURE 20

PSCA Immunostaining of Primary Tumors

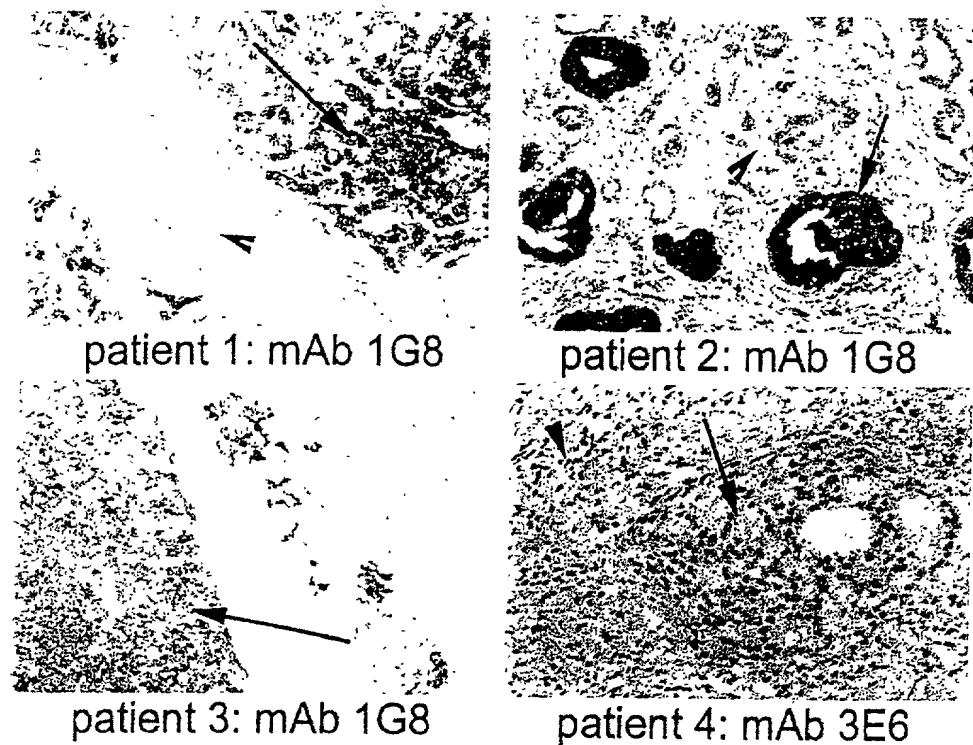


FIGURE 21

44-200-26460



FIGURE 22



FIGURE 23



FIGURE 24

FIGURE 25 - FIGURE 25

1000 ft PSC
1000 ft thick

FIGURE 25

107283 4 2000

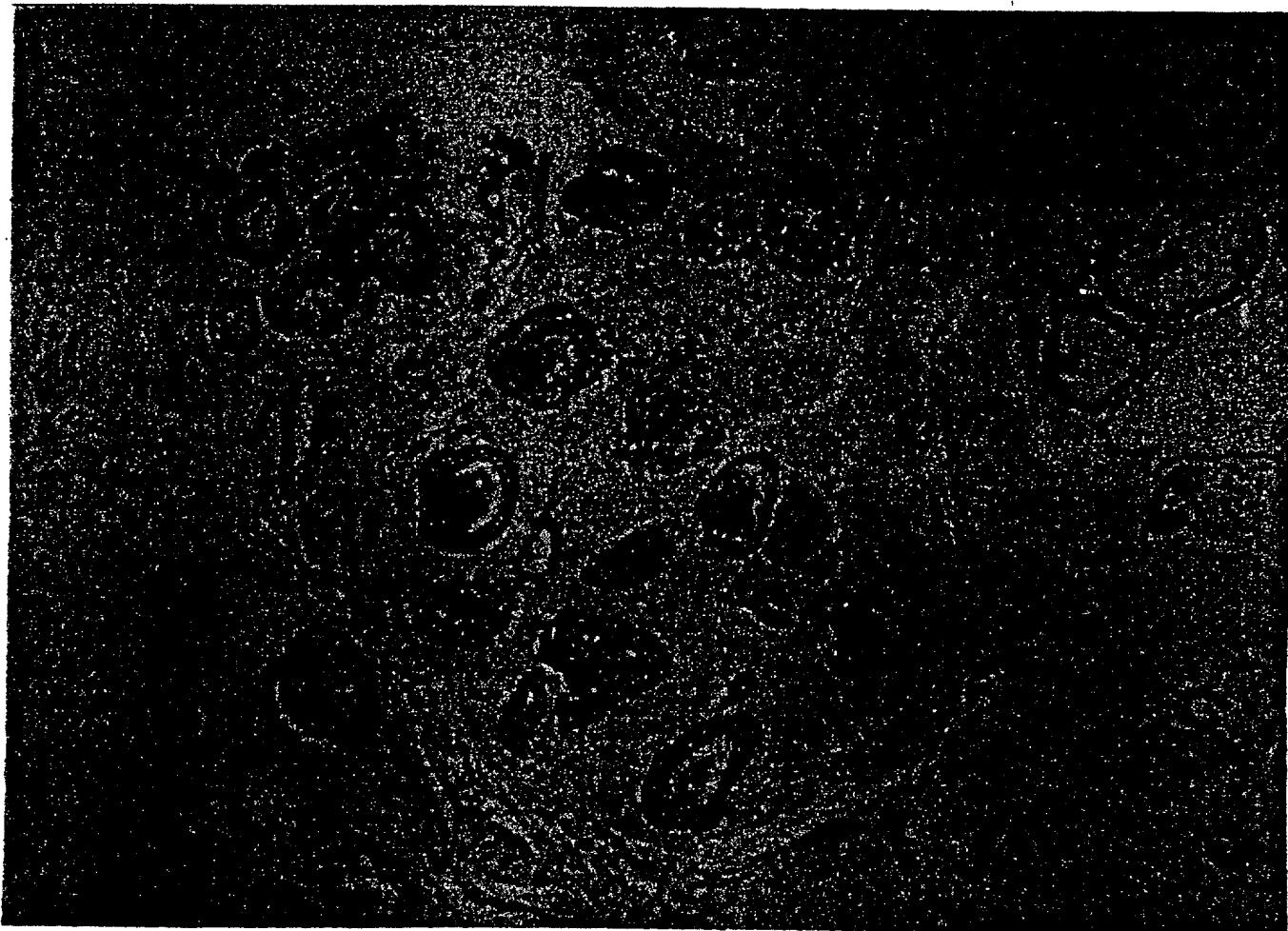


FIGURE 26

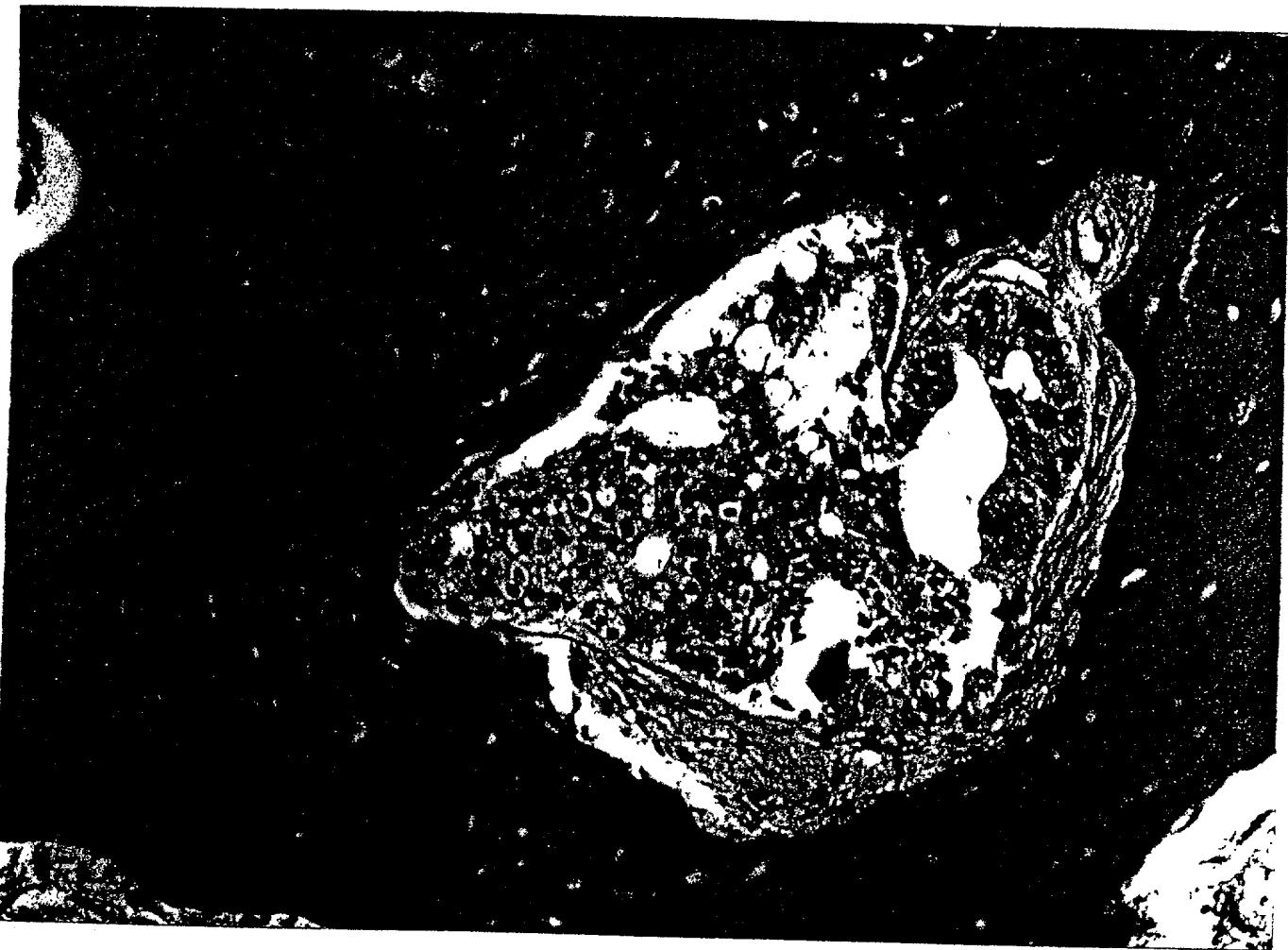
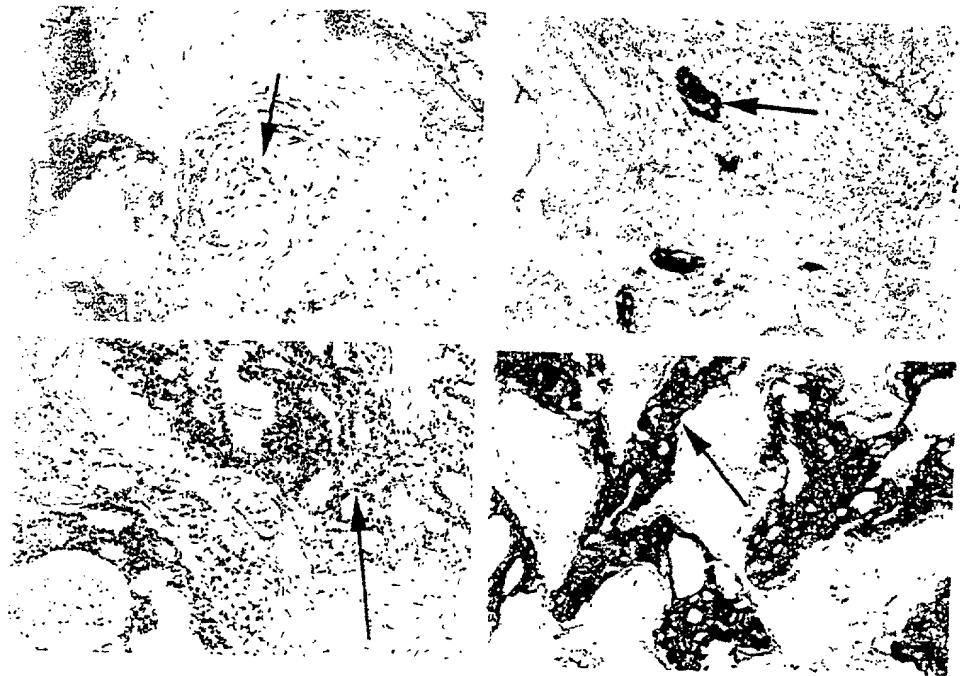


FIGURE 27

PSCA Immunostaining of Bony Metastases



Patient 5: H and E
and mAb 1G8

Patient 4: H and E
and mAb 3E6

FIGURE 28

059214256

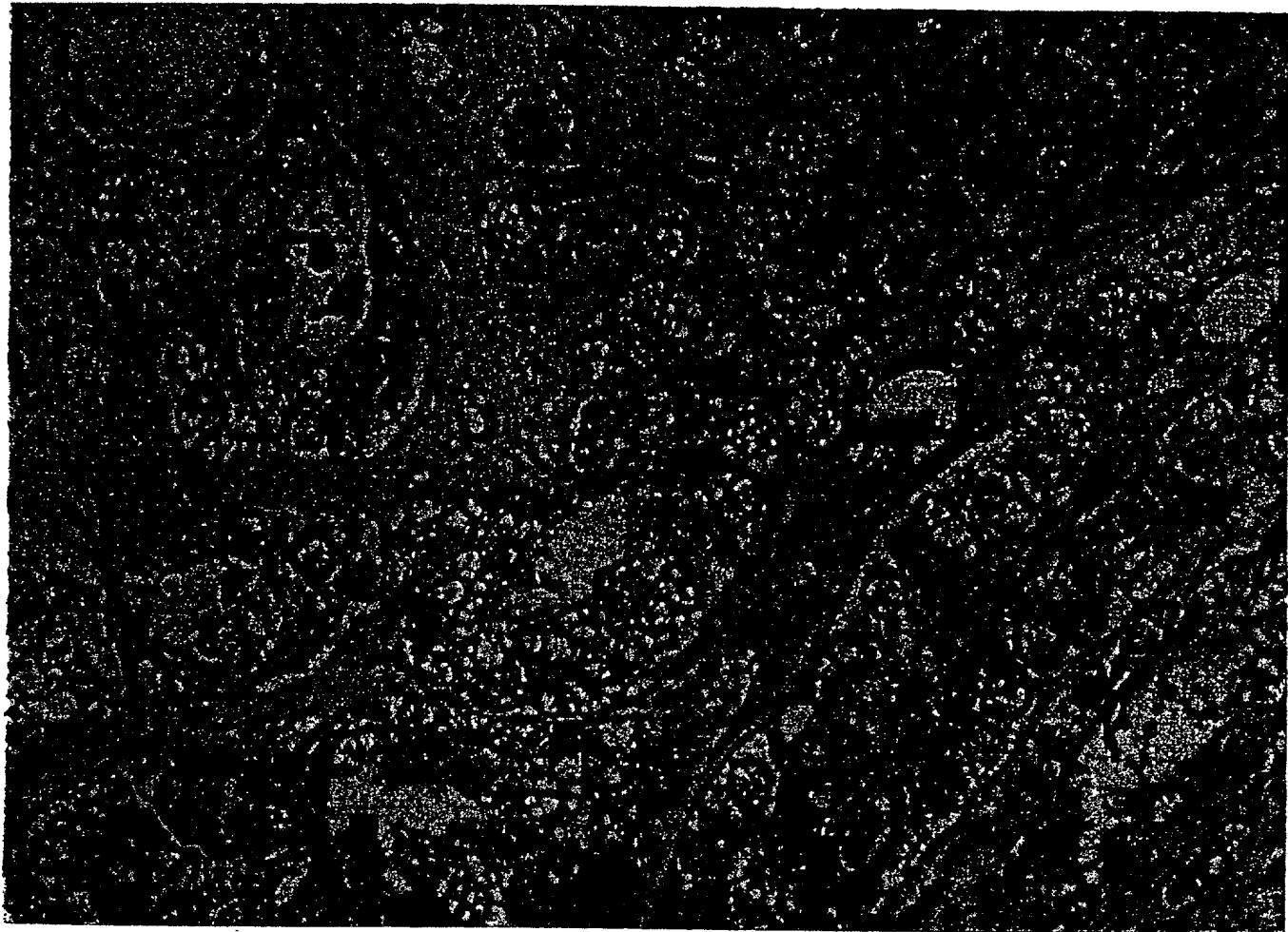


FIGURE 29

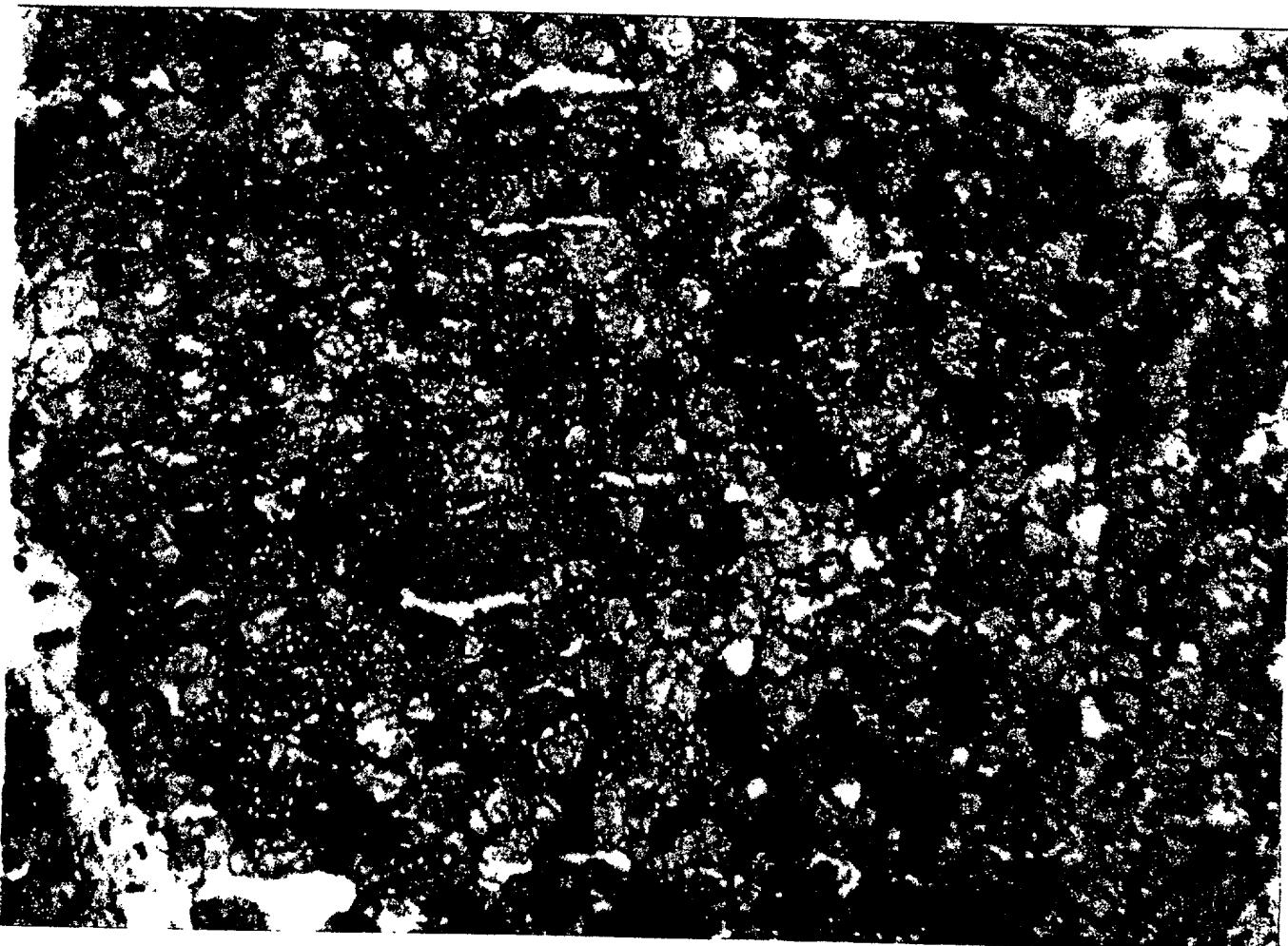


FIGURE 30

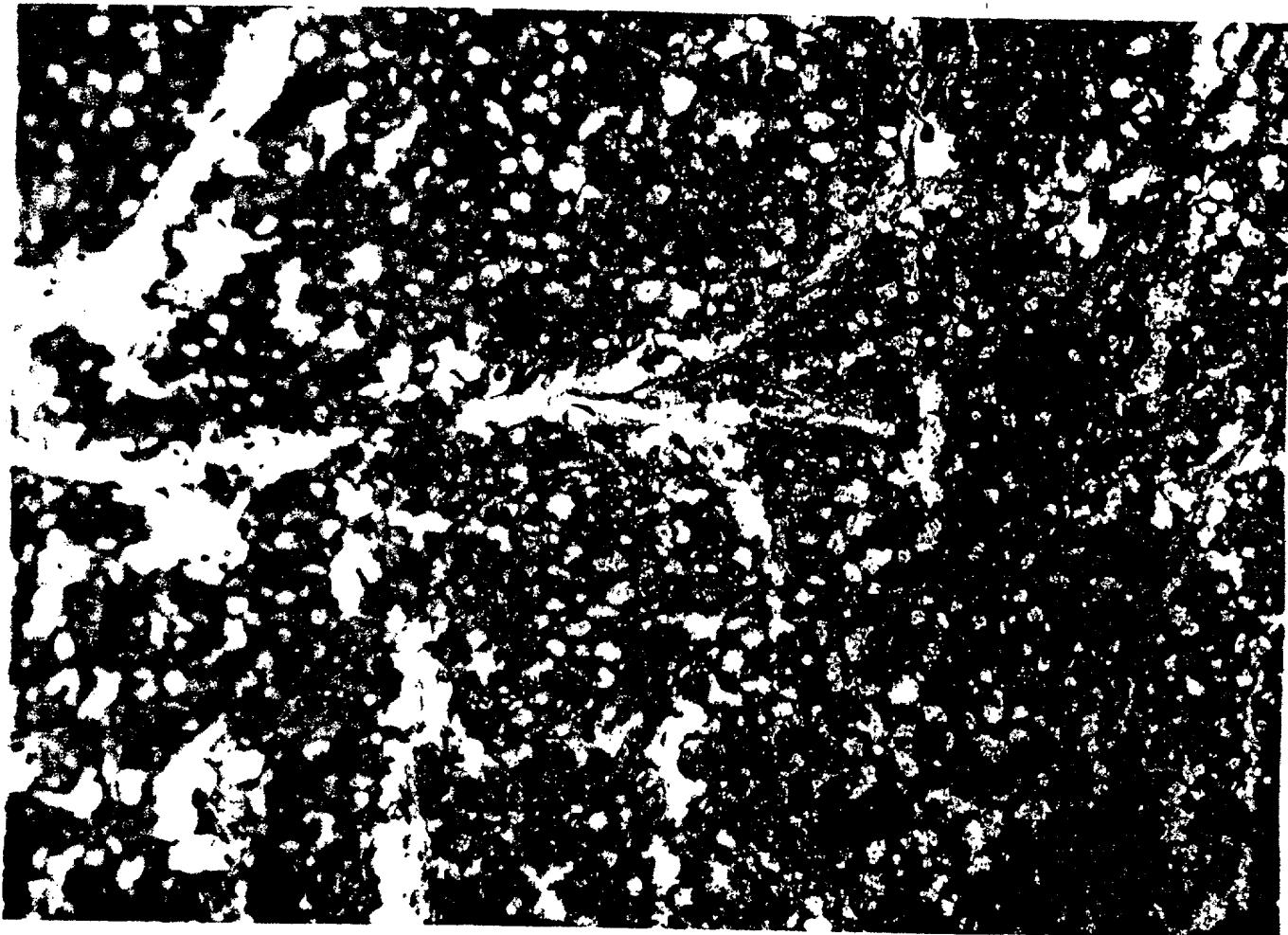


FIGURE 31

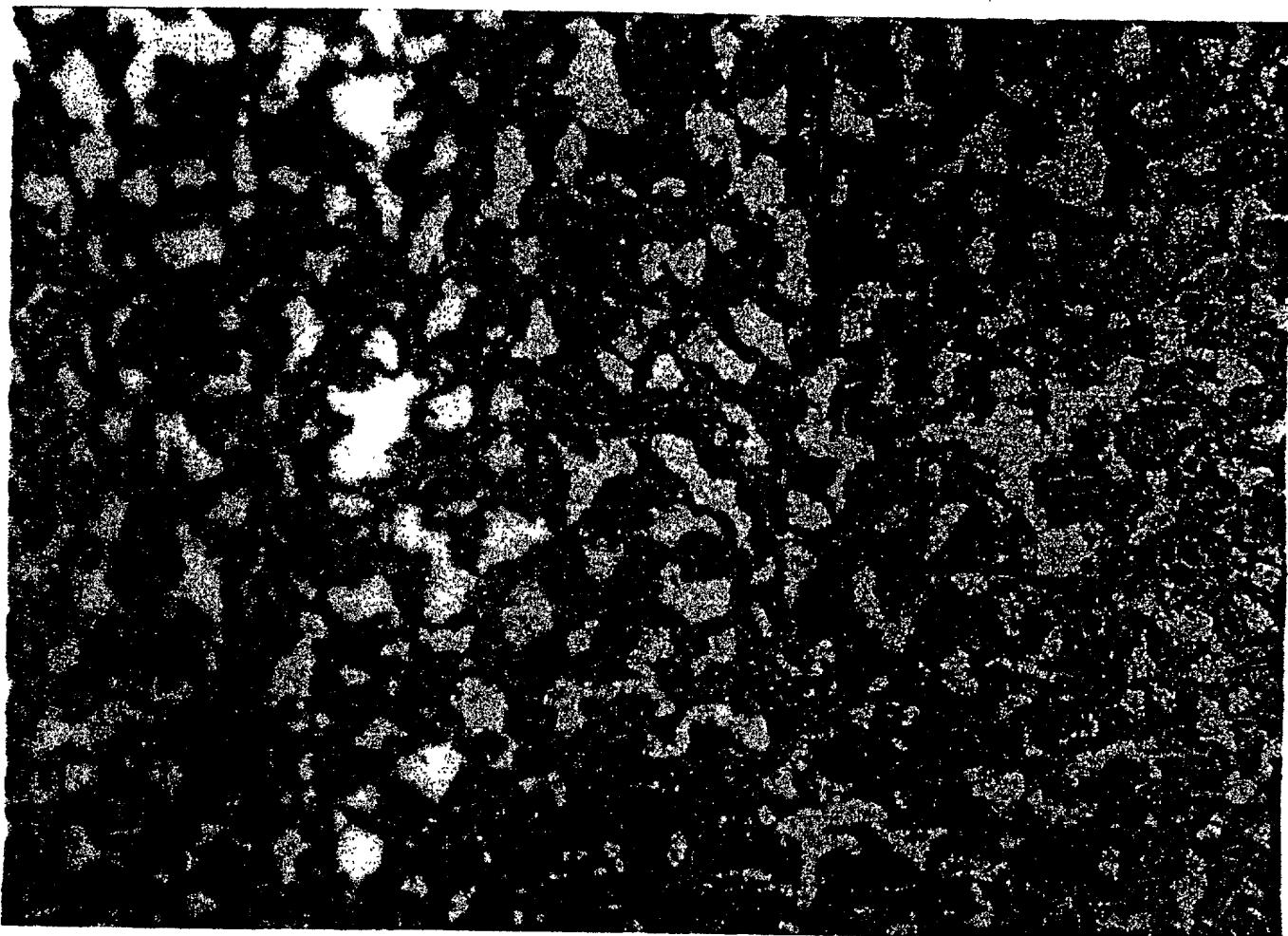


FIGURE 32

PSCA Expression in LAPC-9 Xenograft by FACS

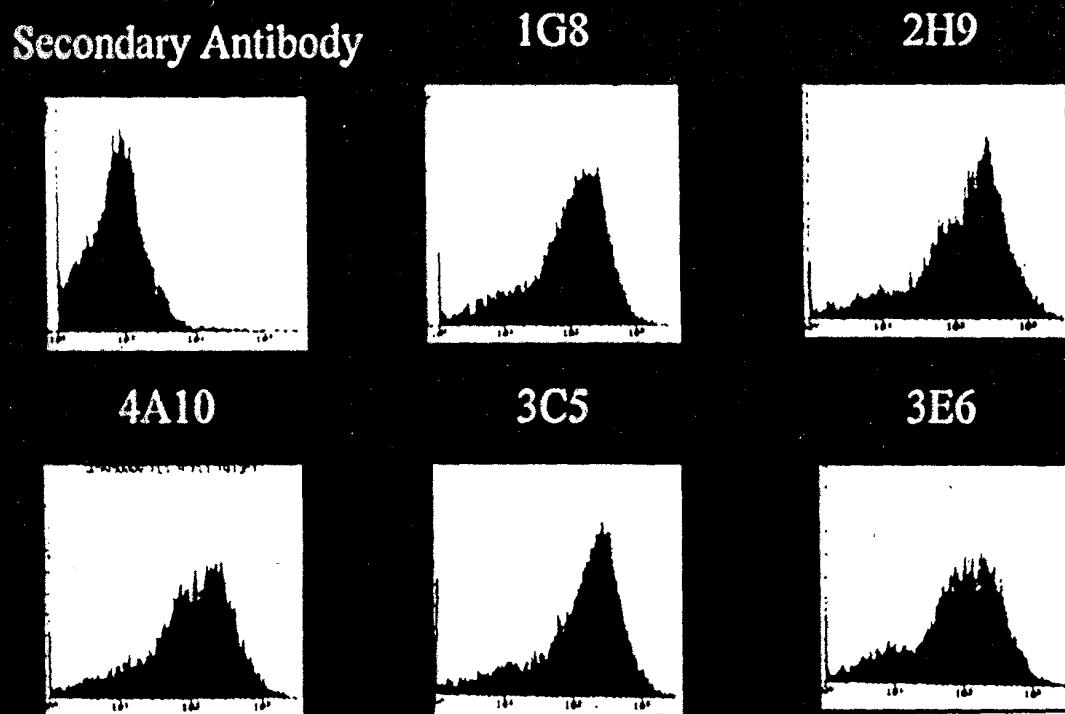


FIGURE 33

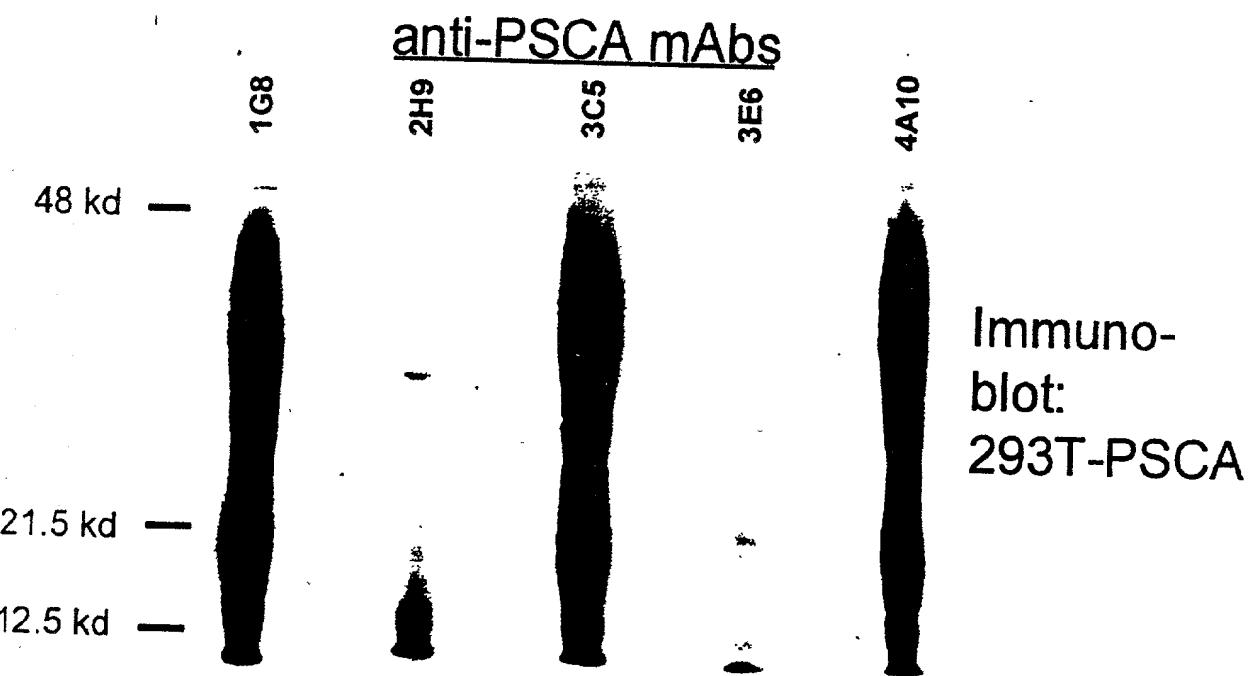


FIGURE 34

Immunofluorescent Staining of LNCaP-PSCA Cells

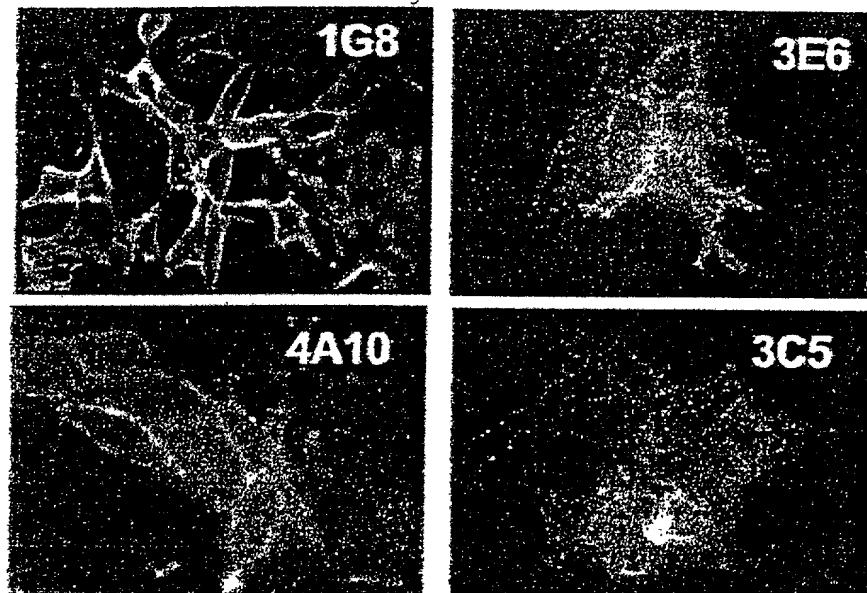


FIGURE 35

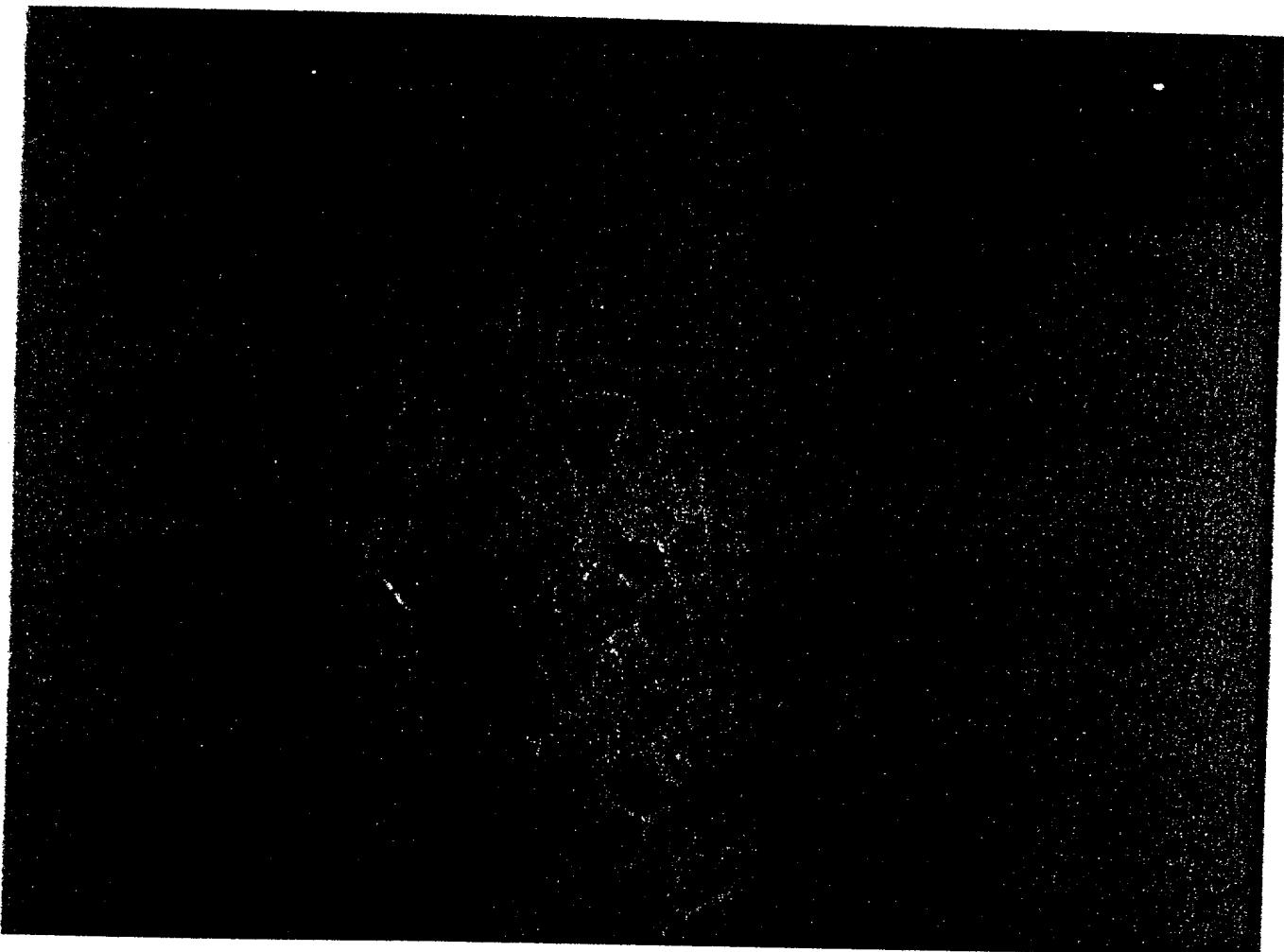


FIGURE 36

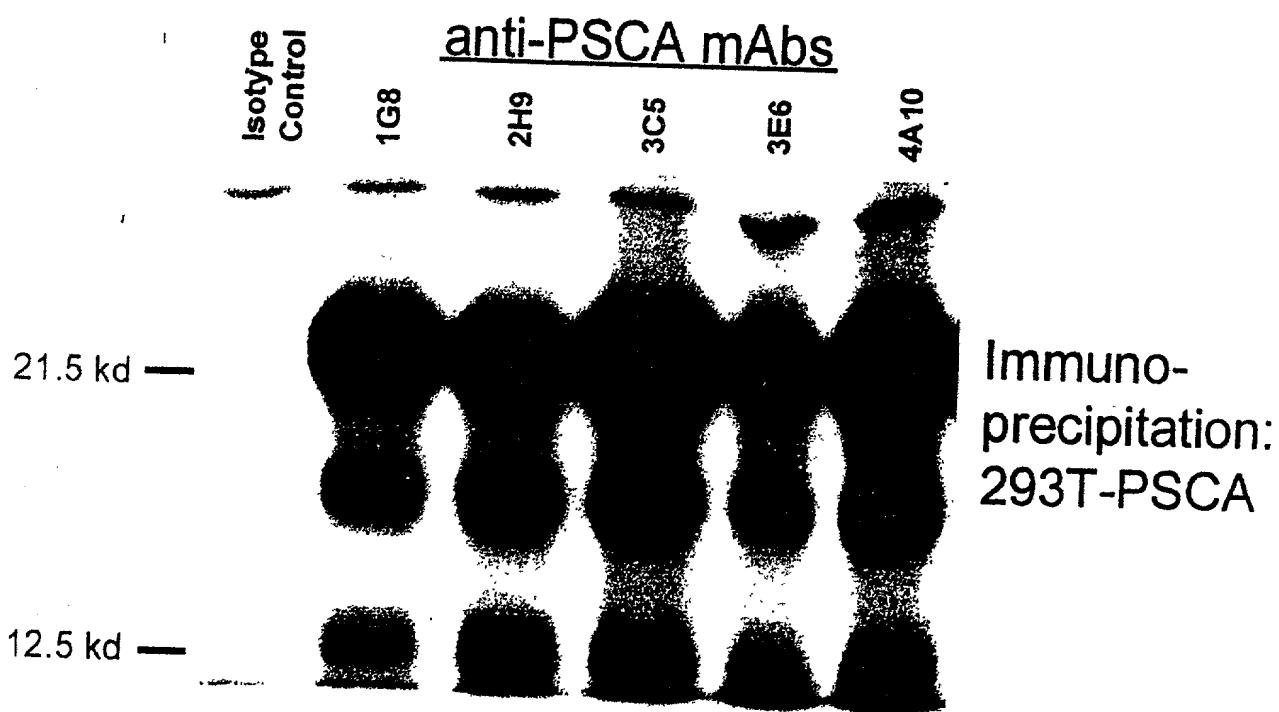


FIGURE 37

Immunohistochemical Staining of Normal Prostate

Normal: Isotype Control



Normal: PSCA mAb 3E6



Normal: PSCA mAb 1G8

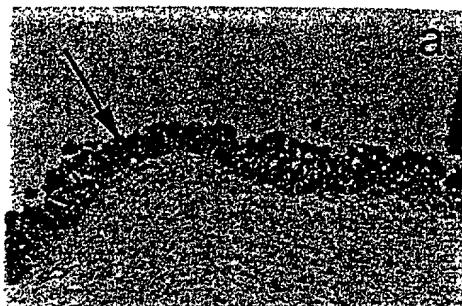


Atrophy: PSCA mAb 2H9

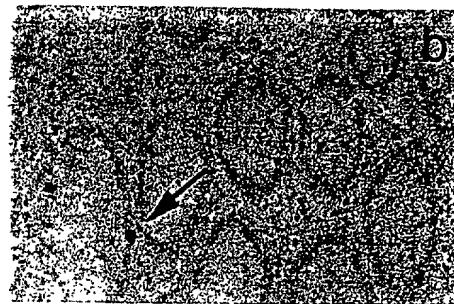


FIGURE 38

A.



Bladder: 1G8



Colon: 1G8

C.



Kidney: 3E6

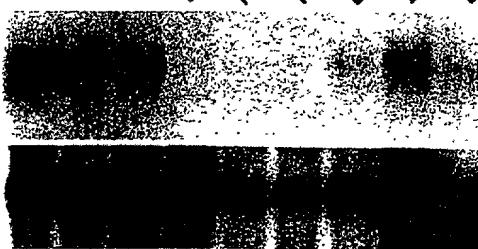
d.



Placenta: 3E6

B.

Prostate
Prostate
Prostate
Kidney
Kidney
Kidney
Bladder
Bladder
Bladder
LAPC9

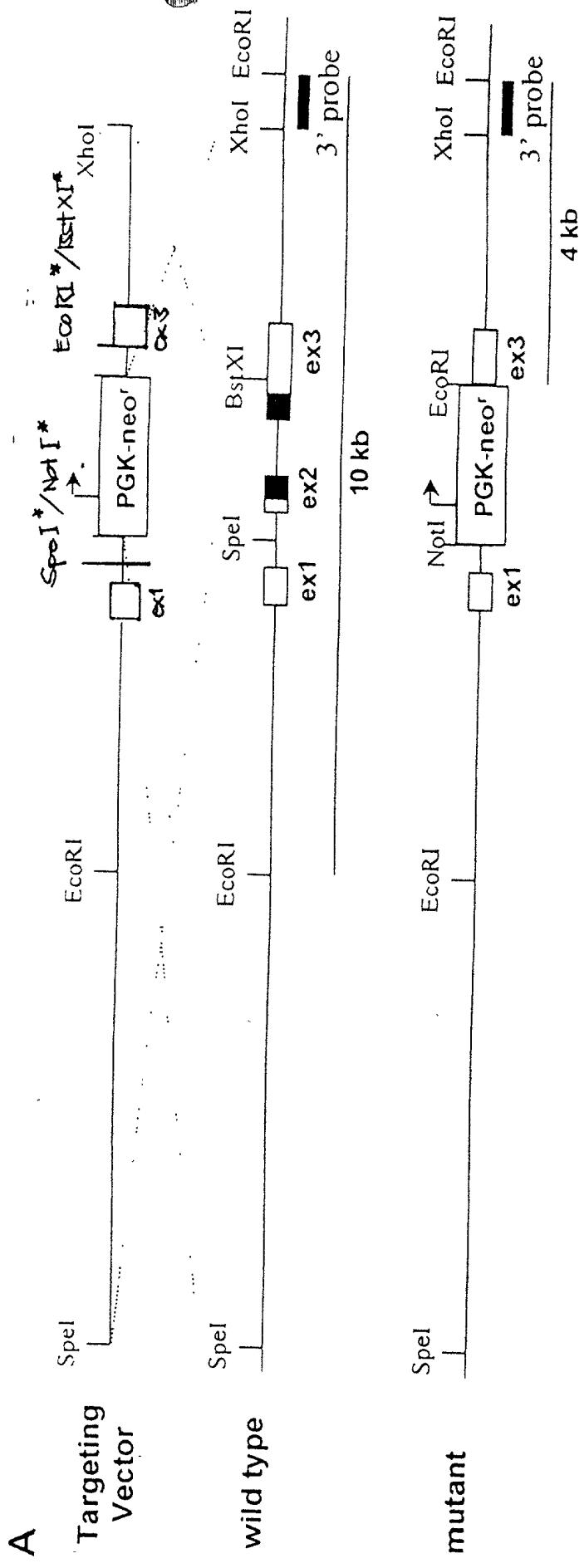


PSCA

Actin

FIGURE 39

Targeting of Mouse PSCA Gene



B. Genomic Southern Analysis of ES Cells

- * ex1, 2, and 3 are the exons of PSCA gene.
- * Black boxes of ex2 and ex3 encode PSCA mature protein sequences.
- * ES genomic DNAs were digested with EcoRI, followed by Southern hybridization using 3' probe.

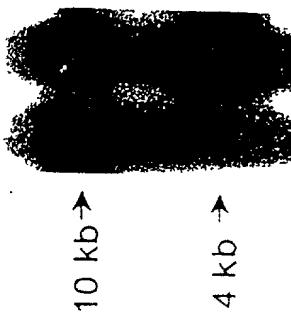
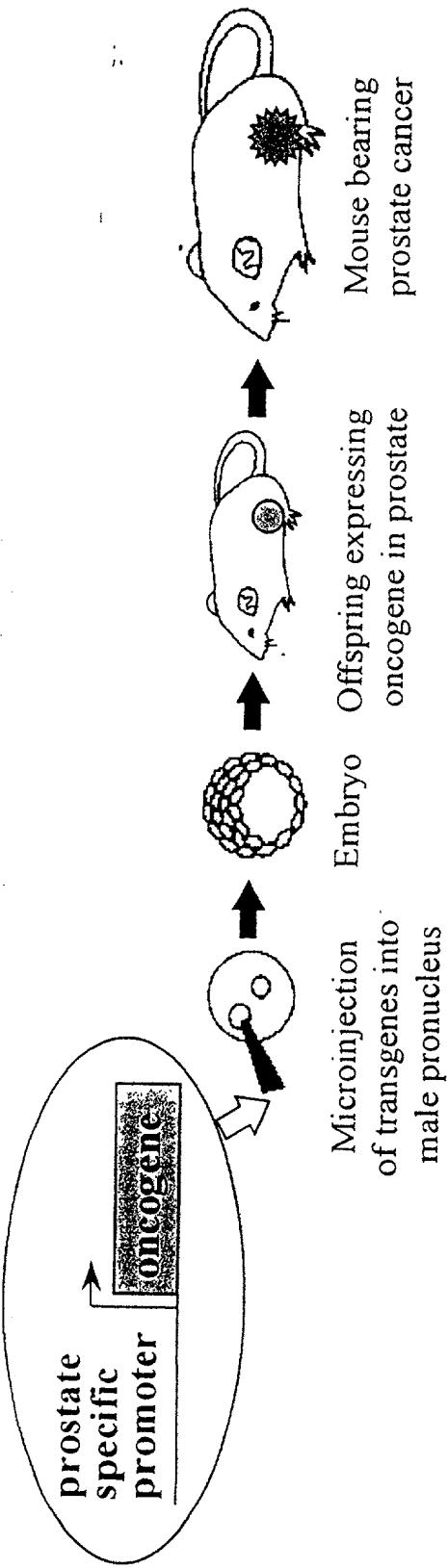


FIGURE 4C

Transgenic Mouse Models of Prostate Cancer



Transgene	Target tissues	Characteristics
C3(1) (-3 kb)/ SV40 large+small T <i>Maroulakou et al.</i> 1994 <i>PNAS</i>	prostate (secretory cells) urethral, mammary and sweat gland	Low-grade PIN 8-12 wks High-grade PIN 8-12 wks Invasive carcinoma 28 wks No metastases
Probasin (-426 bp)/ SV40 large+small T <i>Greenberg et al.</i> 1995 <i>PNAS</i>	prostate (secretory cells)	Low-grade PIN 5-8 wks High-grade PIN 8-12 wks Invasive carcinoma 12 wks Metastases in lymph node, lung, liver and bone
Cryptdin2 (-6.5 kb)/ SV40 large+small T <i>Garabedian et al.</i> 1998 <i>PNAS</i>	prostate (neuroendocrine cells) small intestine	Low-grade PIN 8-12 wks High-grade PIN 8-12 wks Invasive carcinoma 16 wks Metastases in lymph node, lung, liver and bone

FIGURE 41

Reporter Gene Constructs for Transfection Assay

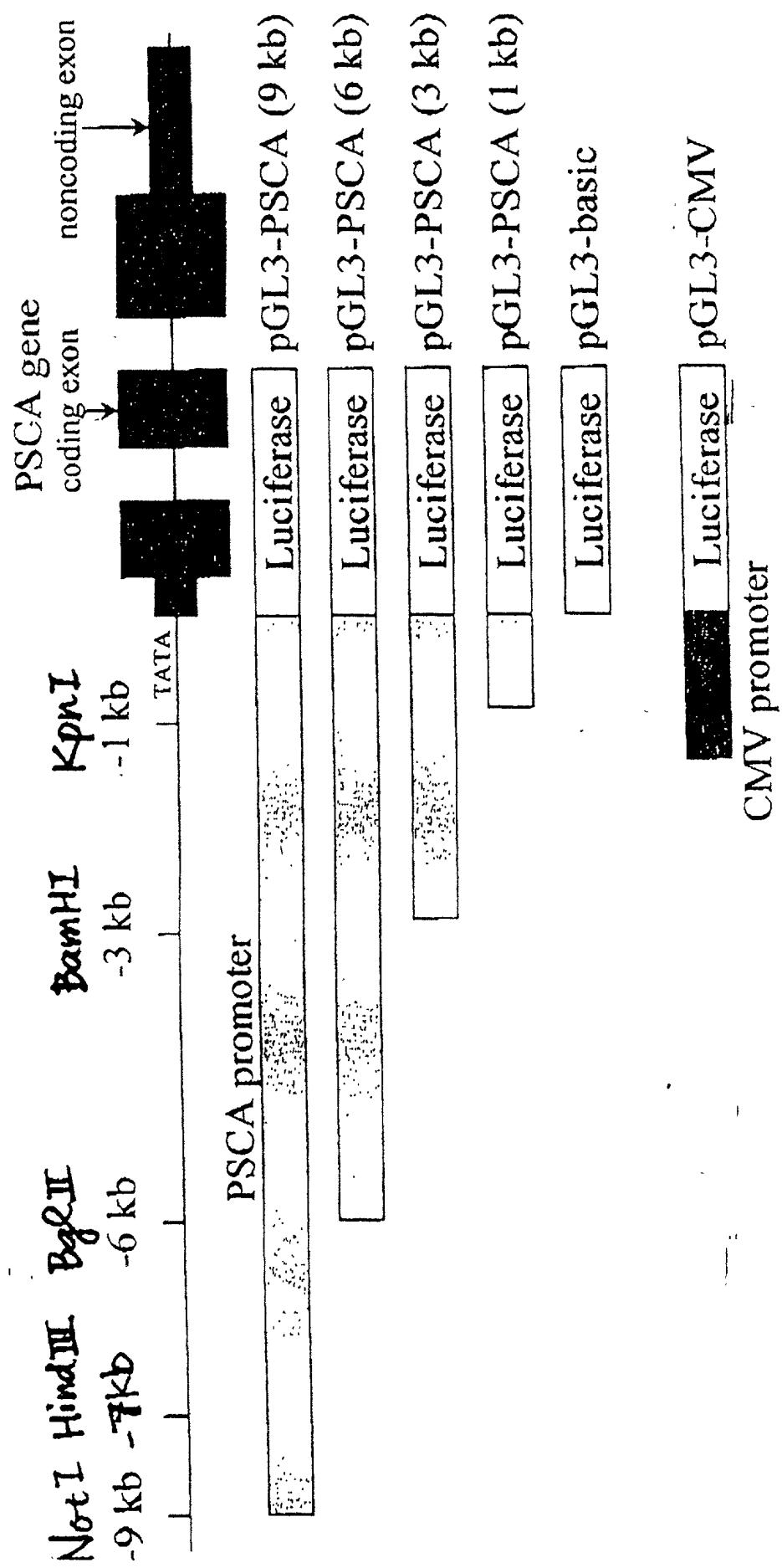


FIGURE 42

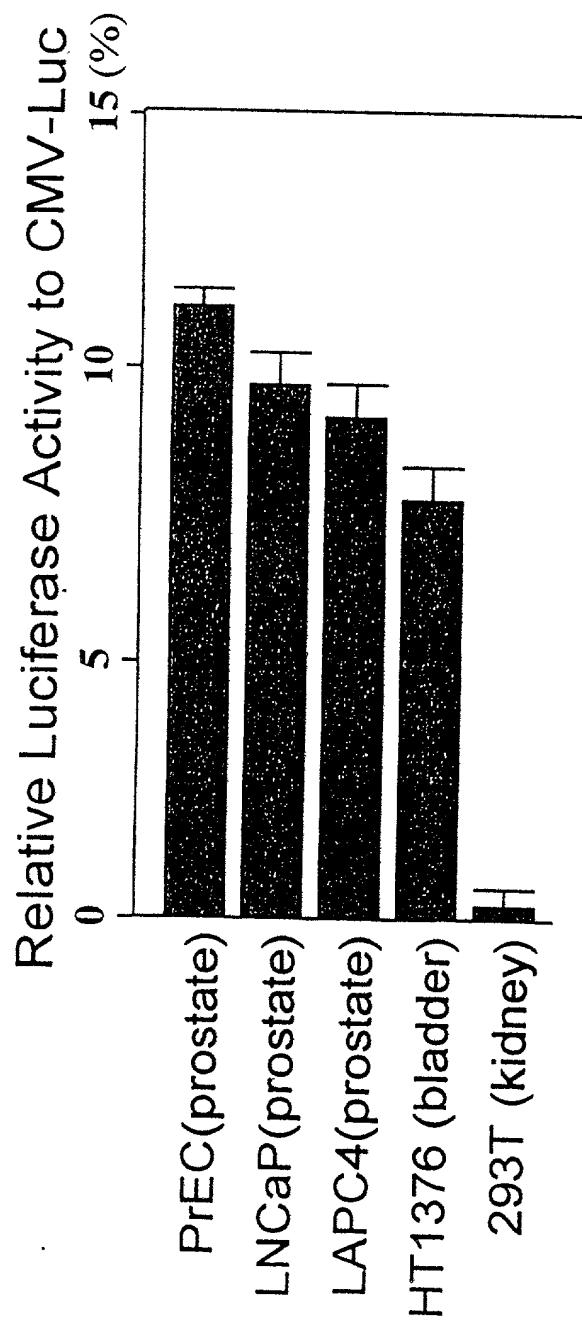


FIGURE 43

Identification of Prostate-Specific Elements Within PSCA Promoter Sequences

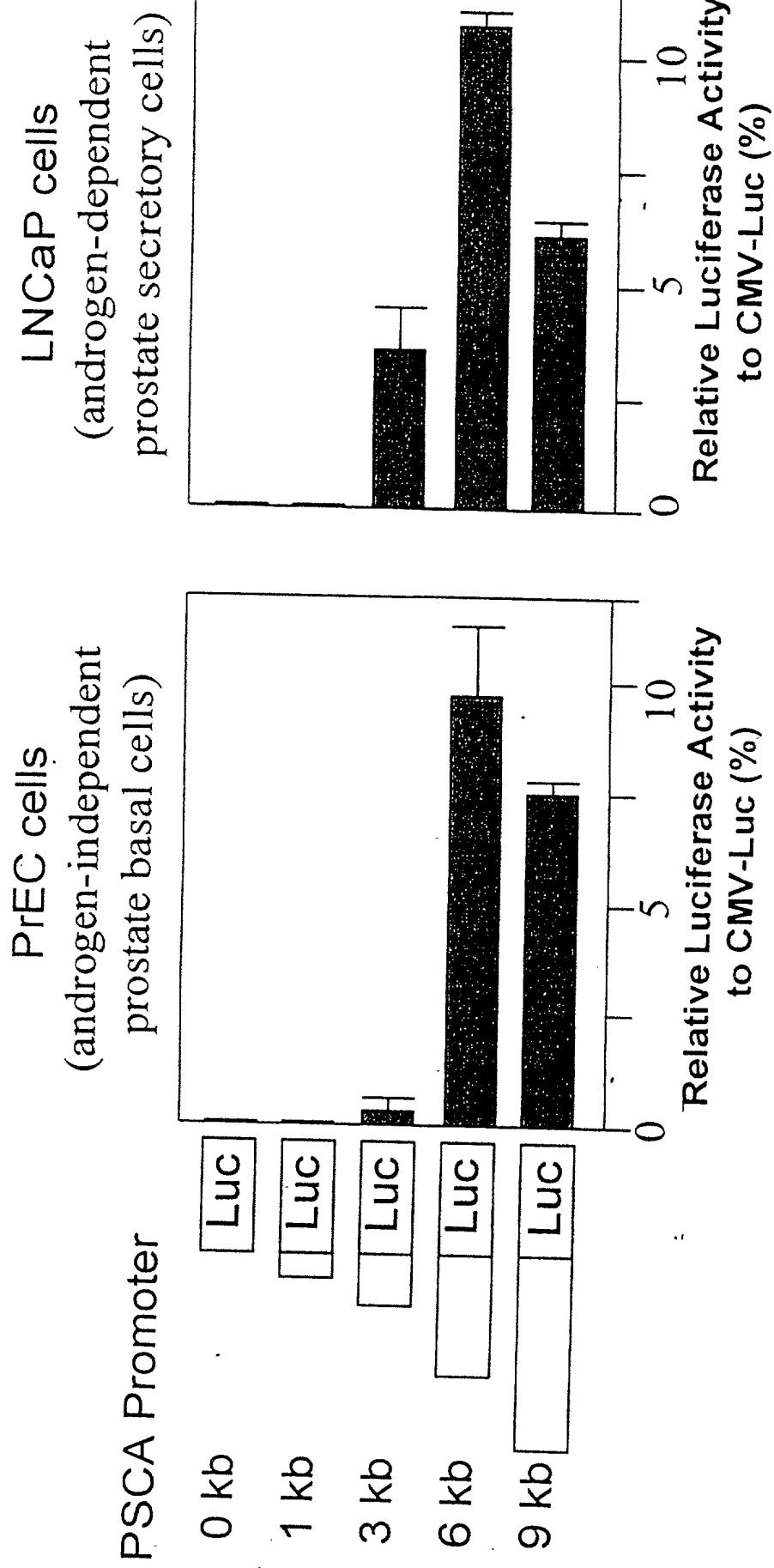


FIGURE 44

Update of Transgenic Mouse Projects

Genomic Structure of PSCA	Number of Founders (DNA positive)
PSCA promoter	2
PSCA(9 kb)-GFP	1
PSCA(6 kb)-GFP	6
PSCA(9 kb)-GFP-3' hGH	8
PSCA(6 kb)-GFP-3' hGH	3
PSCA(9 kb)-SV40TAG	9
PSCA(6 kb)-SV40TAG	3

Diagram illustrating the genomic structures of the PSCA constructs:

- PSCA promoter:** Shows the PSCA promoter region followed by the ATG start site and the first exon (exon 1).
- PSCA(9 kb)-GFP:** Shows the PSCA promoter followed by a 9 kb genomic region containing exon 1, exon 2, and exon 3, which is then fused to the GFP gene.
- PSCA(6 kb)-GFP:** Shows the PSCA promoter followed by a 6 kb genomic region containing exon 1, exon 2, and exon 3, which is then fused to the GFP gene.
- PSCA(9 kb)-GFP-3' hGH:** Shows the PSCA promoter followed by a 9 kb genomic region containing exon 1, exon 2, and exon 3, which is then fused to the GFP gene and includes a 3' hGH polyA signal.
- PSCA(6 kb)-GFP-3' hGH:** Shows the PSCA promoter followed by a 6 kb genomic region containing exon 1, exon 2, and exon 3, which is then fused to the GFP gene and includes a 3' hGH polyA signal.
- PSCA(9 kb)-SV40TAG:** Shows the PSCA promoter followed by a 9 kb genomic region containing exon 1, exon 2, and exon 3, which is then fused to the SV40 polyA signal (TAG).
- PSCA(6 kb)-SV40TAG:** Shows the PSCA promoter followed by a 6 kb genomic region containing exon 1, exon 2, and exon 3, which is then fused to the SV40 polyA signal (TAG).

FIGURE 45

Negative tissues

Stomach
Small intestine
Colon
Seminal Vesicle
Urethra
Testis
Liver
Kidney
Lung
Brain
Heart
Skeletal muscle
Ovary
Uterus

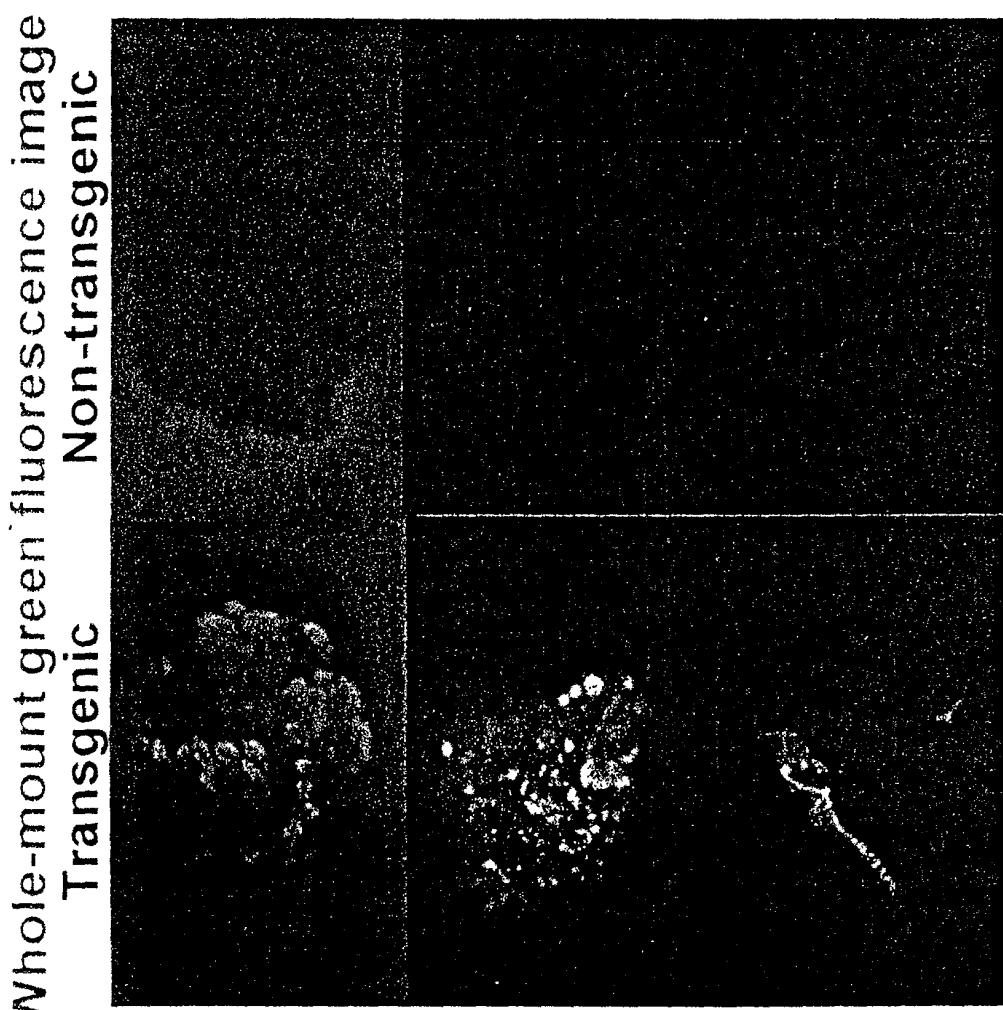


FIGURE 47

RT-PCR



Northern Analysis

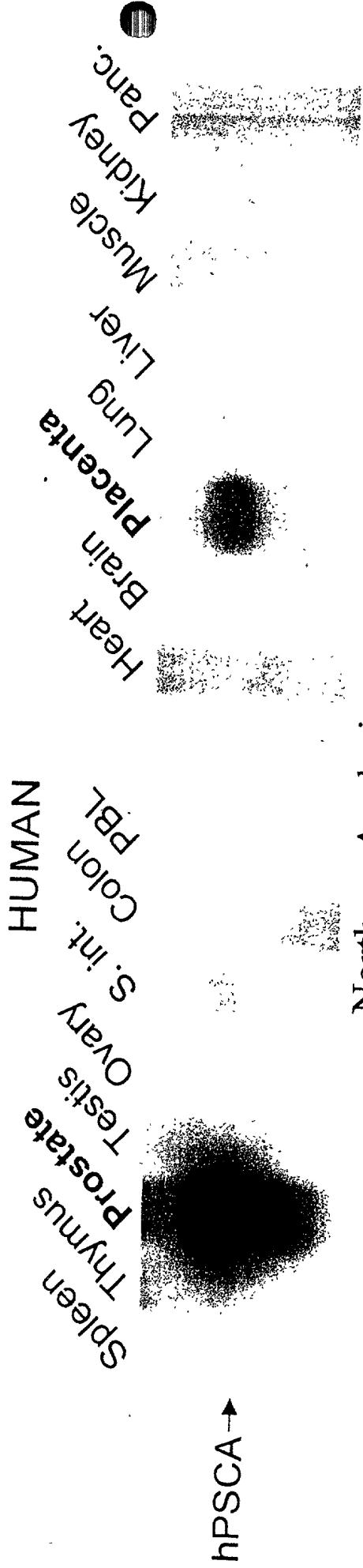
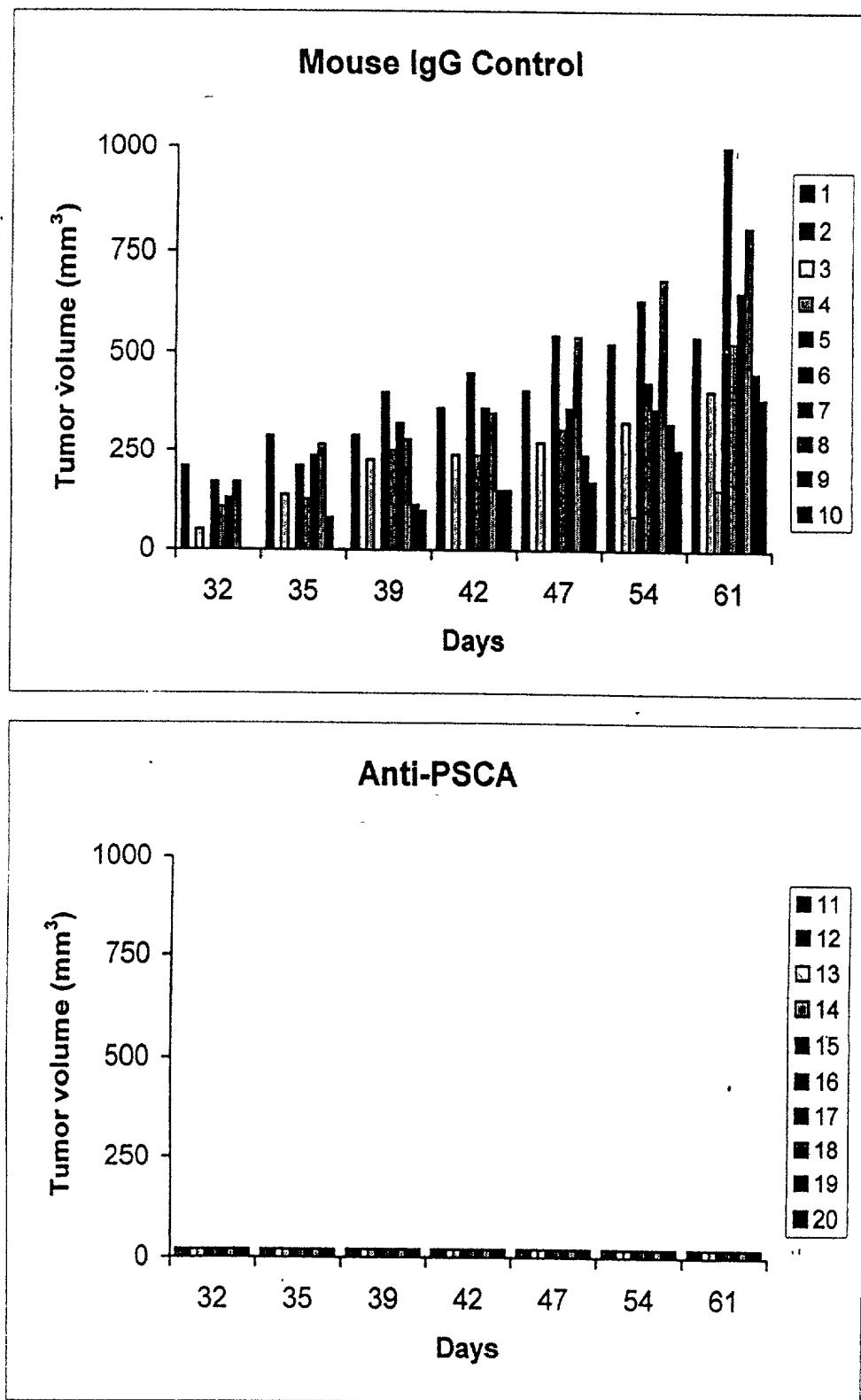


FIG. 48



A**FIG. 49**

<u>mAb</u>	<u>Isotype</u>	<u>F(18-98)</u>	<u>N(2-50)</u>	<u>M(46-109)</u>	<u>C(85-123)</u>
1G8	K	1.485	0.004	1.273	0.003
2A2	K	0.973	0.631	0.023	0.010
2H9	K	1.069	1.026	0.002	0.001
3C5	K	1.916	1.709	0.006	0.002
3E6	K	1.609	0.036	1.133	2.118
3G3	K	2.805	1.731	0.004	0.000
4A10	K	1.053	0.493	0.000	0.001

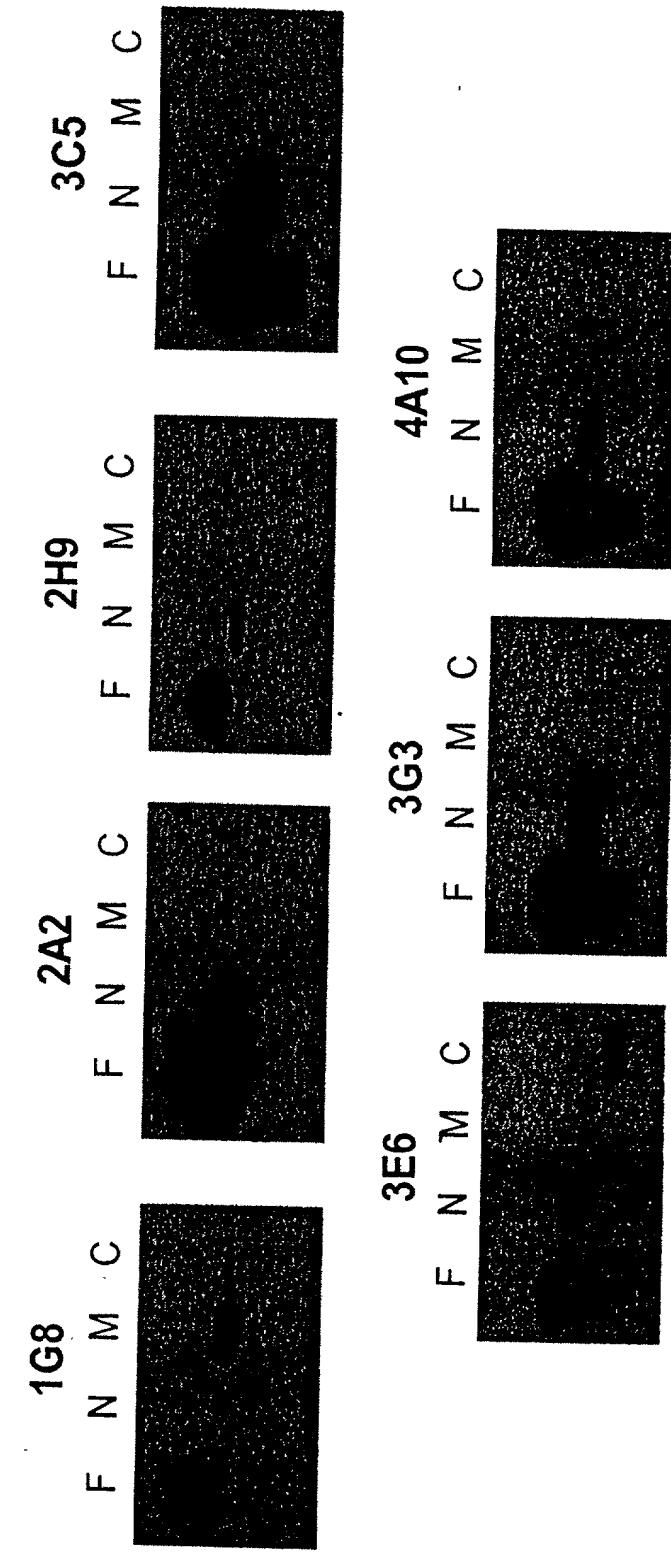
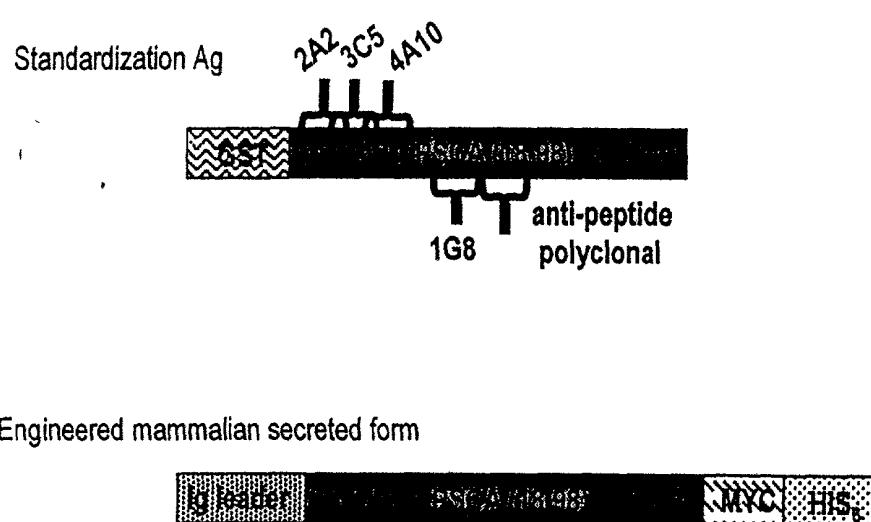
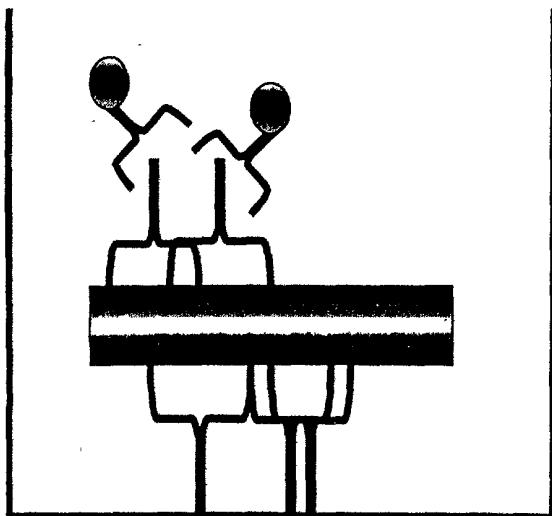


FIG. 50

A



B



Anti-IgG2a HRP

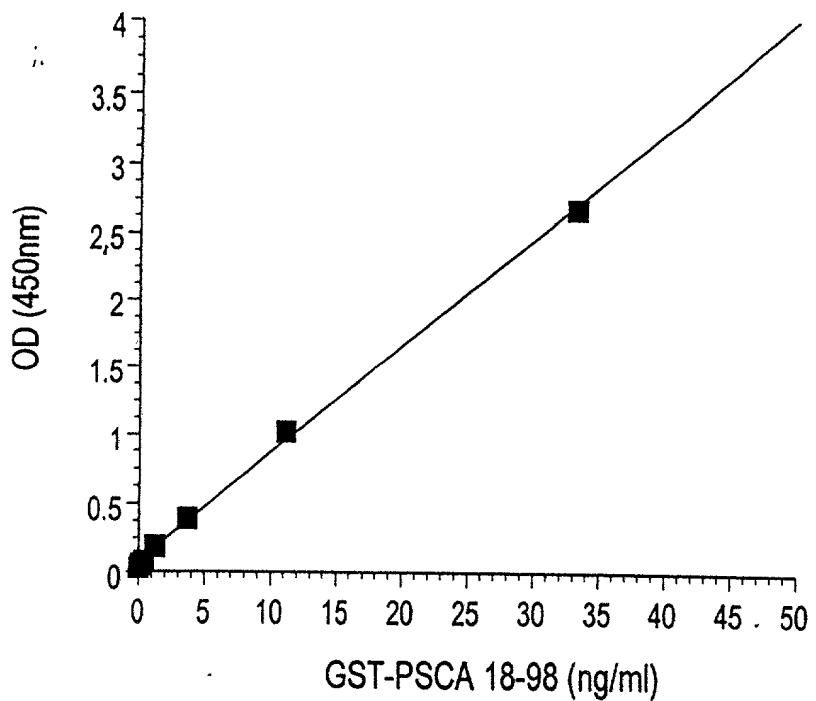
Anti-PSCA mAbs 3C5+4A10+2A2 (IgG2a)

PSCA

Affinity purified anti-peptide polyclonal
+ mAb 1G8 (IgG1)

FIG. 51

A



B

<u>Sample</u>	<u>OD+range (n=2)</u>	<u>ng/ml</u>
vector	0.005+0.001	ND
vector+hu serum	0.004+0.001	ND
secPSCA	2.695+0.031	32.92
secPSCA+hu serum	2.187+0.029	26.55

FIG. 52

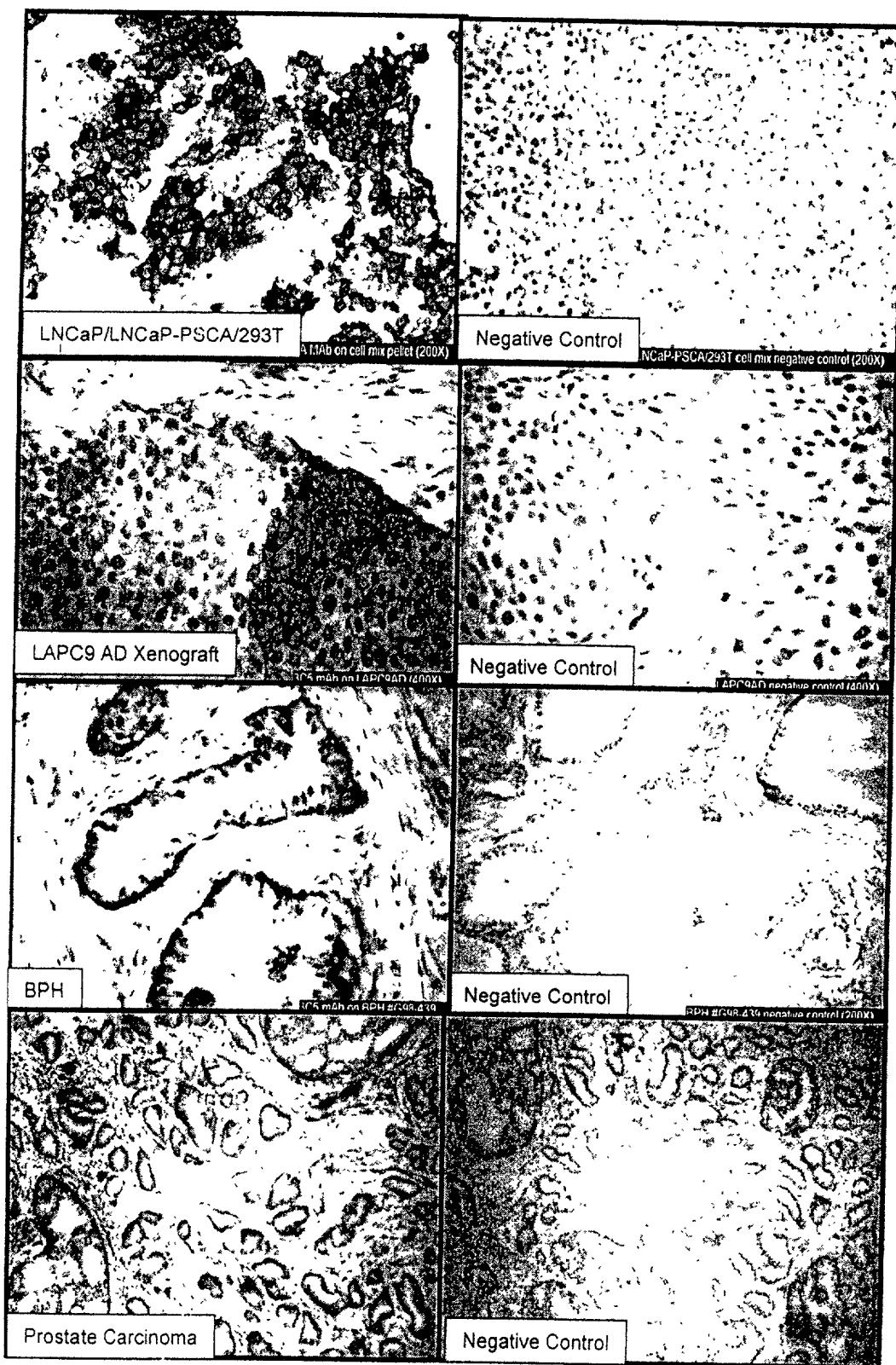


FIG. 53

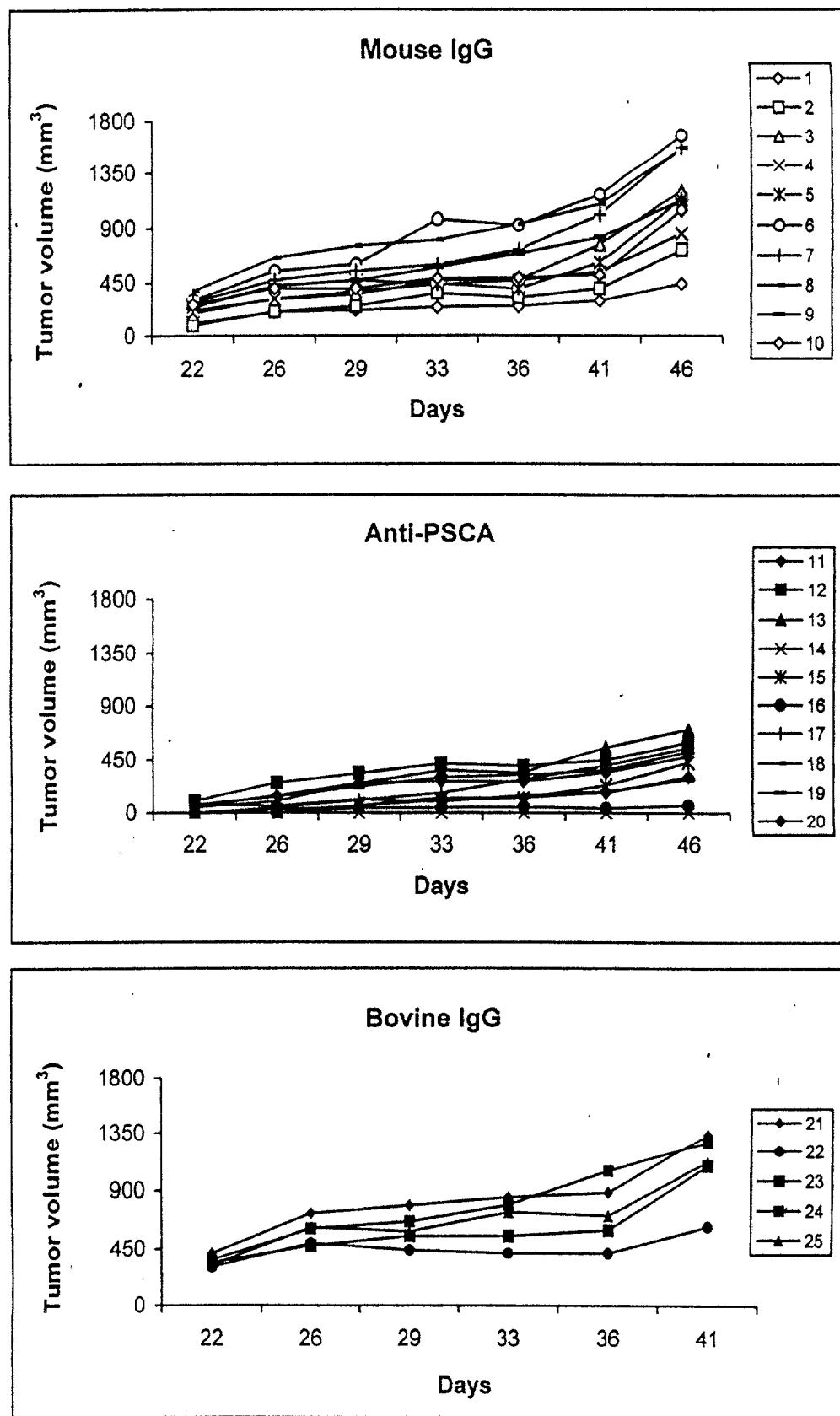


FIG. 54

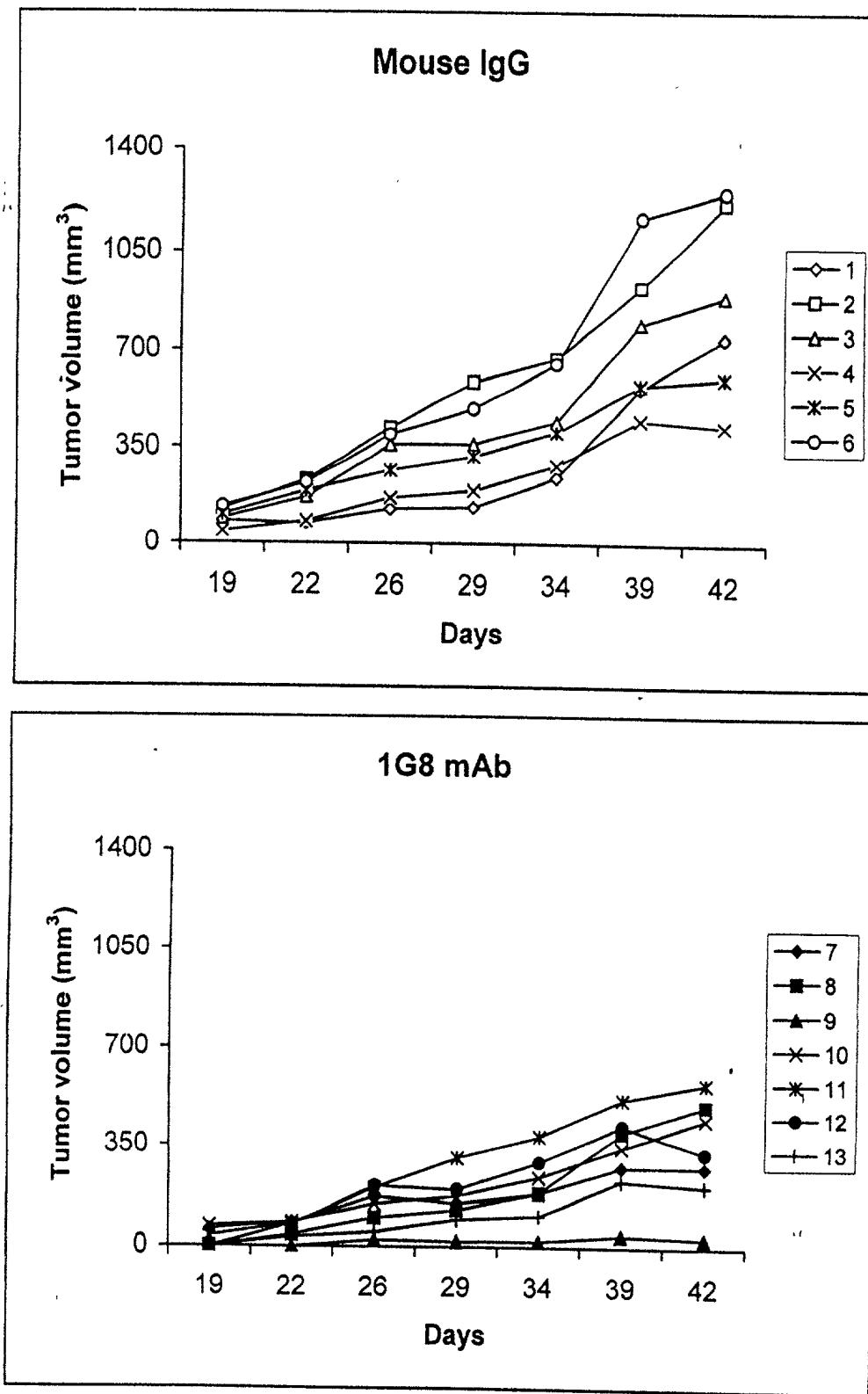


FIG. 55

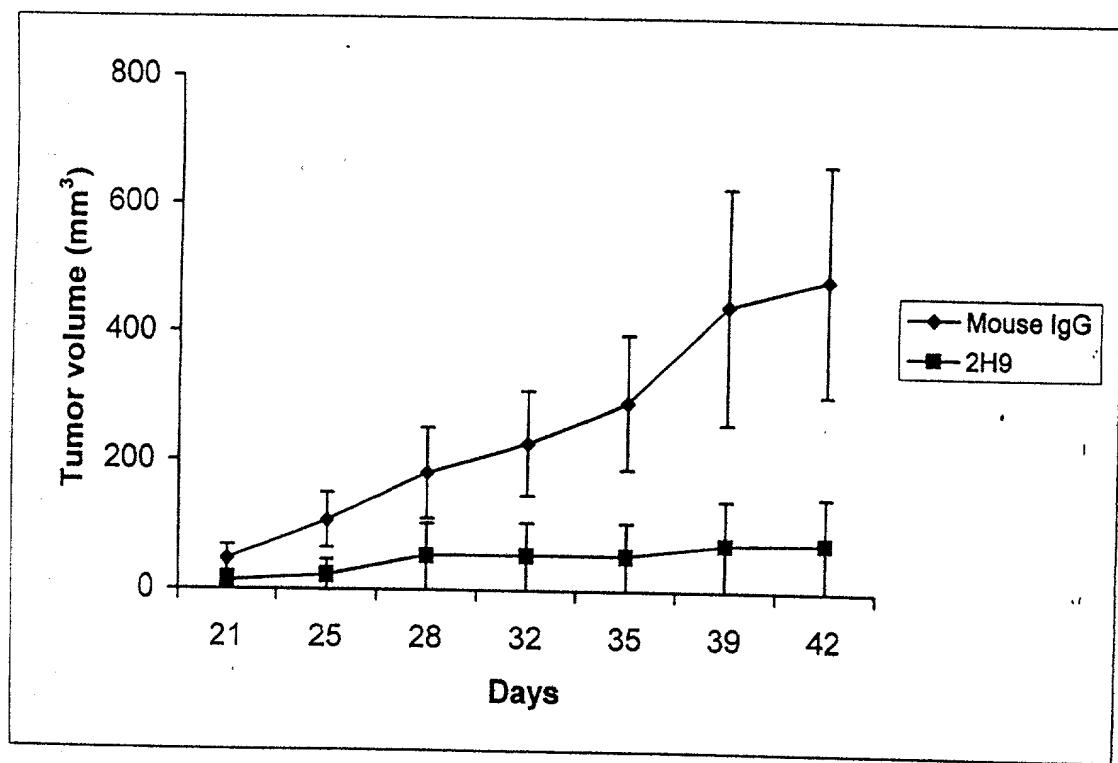
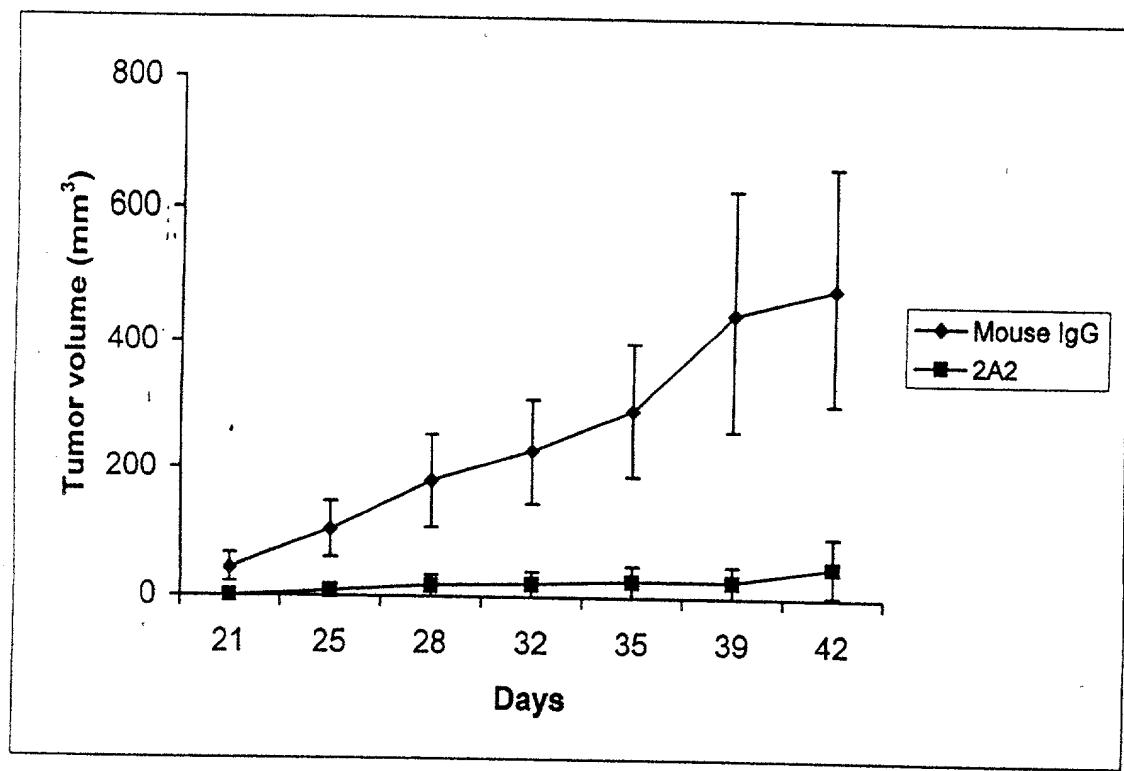


FIG. 56

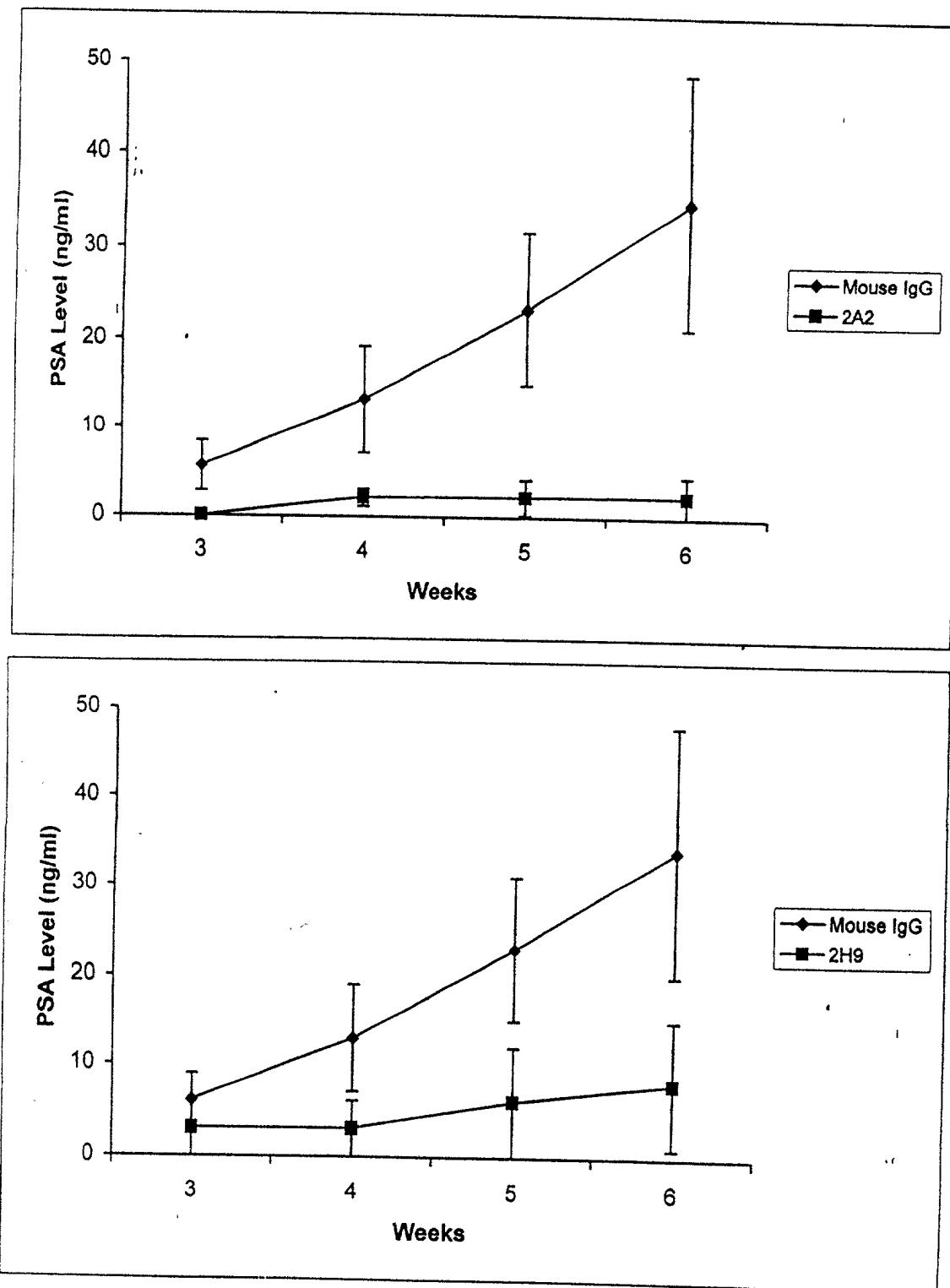


FIG. 57

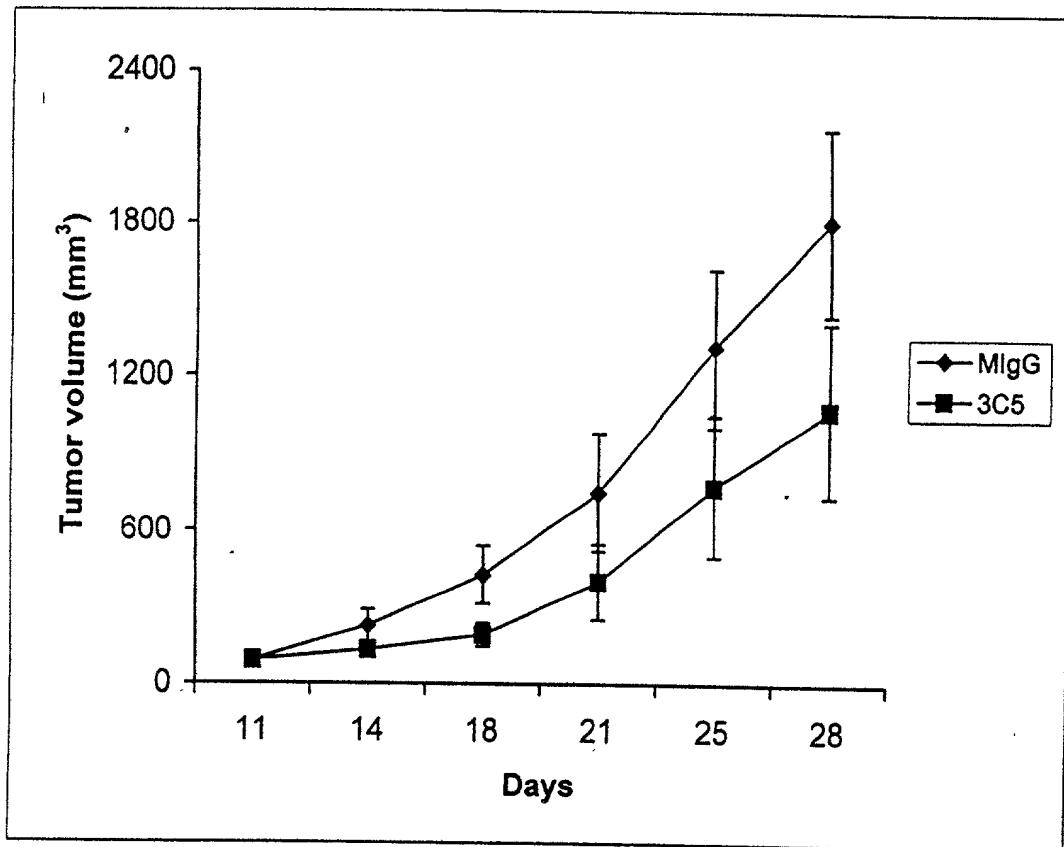


FIG. 58

TGCTTCTCCTGATGGCAGTGGTTAGGAGTCATTAGGAGTTAGCTGCAGCAGTCT 60
C F F L M A V V I G V N S E V Q L Q Q S 20

GGGGCAGAACATTGTGAGGTCAGGGCCTCAGTCAGTTGTCCTGCACAGCTCTGGCTTC 120
G A E L V R S G A S V K L S C T A S G F 40

CDR1
AACATTAAGACTACTATACACTGGGTGAATCAGAGGCCTGACCAGGCCTGGAGTGG 180
N I K D Y Y I H W V N Q R P D Q G L E W 60

CDR2
ATTGGATGGATTGATCCTGAGAATGGTGACACTGAATTGTCCTGAAGTTCCAGGGCAAG 240
I G W I D P E N G D T E F V P K F O G K 80

GCCACTATGACTGCAGACATTTCTCCAACACAGCCTACCTGCACCTCAGCAGCCTGACA 300
A T M T A D I F S N T A Y L H L S S L T 100

CDR3
TCTGAAGACACTGCCGTCTATTACTGTAAAACGGGGGTTCTGGGGCCAAGGGACTCTG 360
S E D T A V Y Y C K T G G F W G Q G T L 120

GTCACTGTCTCTGCAGCCAAACGACACCCCCATCTGTCTATCCACTG
V T V S A A K T T P P S V Y P L

FIG. 59

TTGGTAGAACAGCCTCAGATGTCCACTCCCAGGTCCAAC TG CAG CAAC CTGGTCTGAA 60
L V A T A S D V H S Q V Q L Q Q P G S E 20

CTGGTAGGGCCTGGAACCTCAGTGAAGCTGTCCTGCAAGGCTTCTGGCTATACATTCTCC 120
L V R P G T S V K L S C K A S G Y T F S 40
CDR1

AGCTACTGGATGCAC TGGTGAAGCAGAGGCCTGGACAAGGCCTTGAGTGGATTGGAAAT 180
S Y W M H W V K Q R P G Q G L E W I G N 60

ATTGACCTGGTAGTGGTTACACTAACTACGCTGAGAACCTCAAGACCAAGGCCACACTG 240
I D P G S G Y T N Y A E N L K T K A T L 80
CDR2

ACTGTAGACACATCCTCCAGCACAGCCTACATGCAGCTCAGCAGCCTGACATCTGAGGAC 300
T V D T S S S T A Y M Q L S S L T S E D 100

TCTGCAGTCTATTACTGTACAAGCCGATCTACTATGATTACGACGGGATTGCTTACTGG 360
S A V Y Y C T S R S T M I T T G F A Y W 120
CDR3

GGCCAAGGGACTCTGGTCACTGTCTGCAGCTACAACAAACAGCCCCATCTGTCTATCCA 420
G Q G T L V T V S A A T T T A P S V Y P 160

CTGGCC
L A

FIG. 60

AATGACTTCGGGTTGAGCTGGGTTTTATTATTGTTCTTTAAAAGGGTCCGGAGTGAA 60
N D F G L S W V F I I V L L K G V R S E 20

GTGAGGCTTGAGGAGTCTGGAGGAGGCTGGGTGCAACCTGGAGGATCCATGAAACTCTCC 120
V R L E E S G G G W V Q P G G S M K L S 40

TGTGTAGCCTCTGGATTTACTTCAGTAATTACTGGATGACTGGGTCCGCCAGTCTCCA 180
C V A S G F T F S N Y W M T W V R Q S P 60
CDR1

GAGAAGGGCTTGAGTGGGTTGCTGAAATTGATTGAGATCTGAAAATTATGCAACACAT 240
E K G L E W V A E I R L R S E N Y A T H 80
CDR2

TATGCGGAGTCTGTGAAAGGAAATTCAACCCTCAAGAGATGATTCCAGAAGTCGTCTC 300
Y A E S V K G K F T I S R D D S R S R L 100

TACCTGCAAATGAACAACTTAAGACCTGAAGACAGTGGATTATTACTGTACAGATGGT 360
Y L Q M N N L R P E D S G I Y Y C T D G 120

CTGGGACGACCTAACTGGGCCAAGGACTCTGGTCACTGTCTCTGCAGCCAAACGACA 420
L G R P N W G Q G T L V T V S A A K T T 140
CDR3

CCCCCATCTGTCTATCCACTGGCCCCTTGTGTA
P P S V Y P L A P C V

FIG. 61

CDR1 Comparisons

1G8	1gG _{1k}	Middle	G	F	N	I	K	D	Y	Y	I	H
2H9	1gG _{1k}	N-Term.	G	F	T	F	S	N	Y	W	M	T
4A10	1gG _{2ak}	N-Term.	G	Y	T	F	S	S	Y	W	M	H

CDR2 Comparisons

1G8	1gG _{1k}	W	I	D	P	E	N	G	D	T	E	F	V	P	K	F	Q	G		
2H9	1gG _{1k}	E	I	R	L	R	S	E	N	Y	A	T	H	Y	A	E	S	V	K	G
4A10	1gG _{2ak}	N	I	D	P	G	S	G	Y	T	N			Y	A	E	N	L	K	T

CDR3 Comparisons

1G8	1gG _{1k}	G	G	F								
2H9	1gG _{1k}	L	G	R	P	N						
4A10	1gG _{2ak}	R	S	T	M	I	T	T	G	F	A	Y

FIG. 62

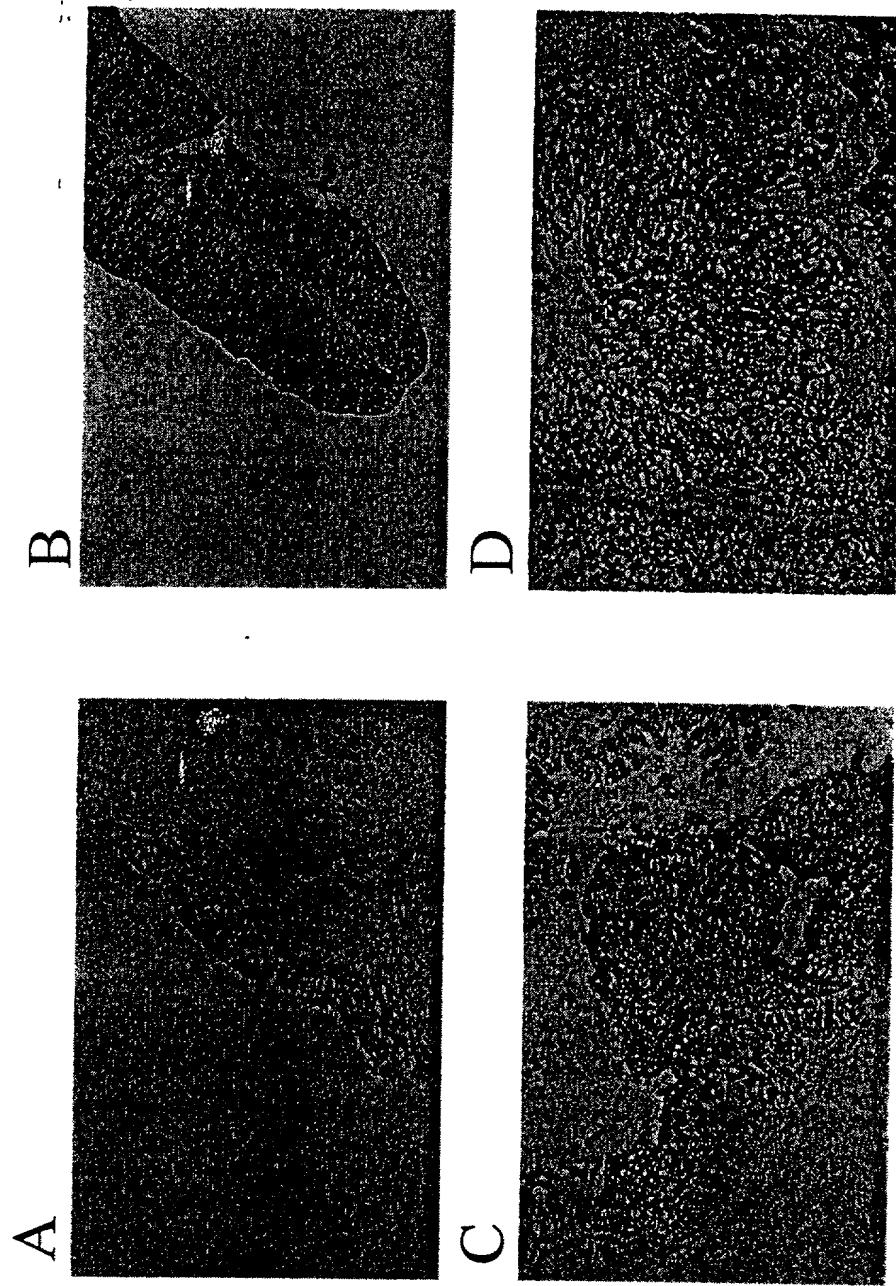


FIG. 63

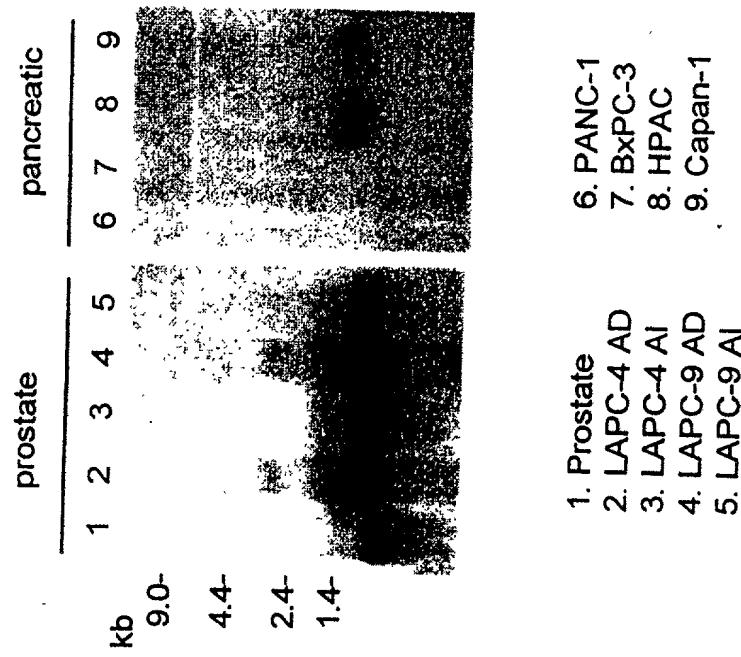
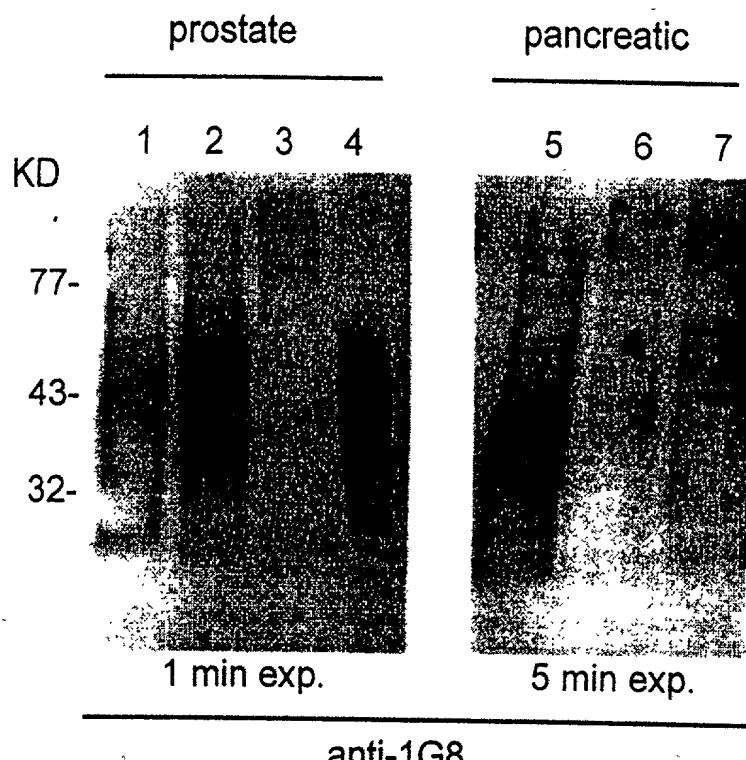


FIG. 64



anti-1G8

1. LAPC-4 AD
2. LAPC-9 AI
3. LNCaP
4. LNCaP-PSCA

5. HPAC
6. Capan-1
7. ASPC-1

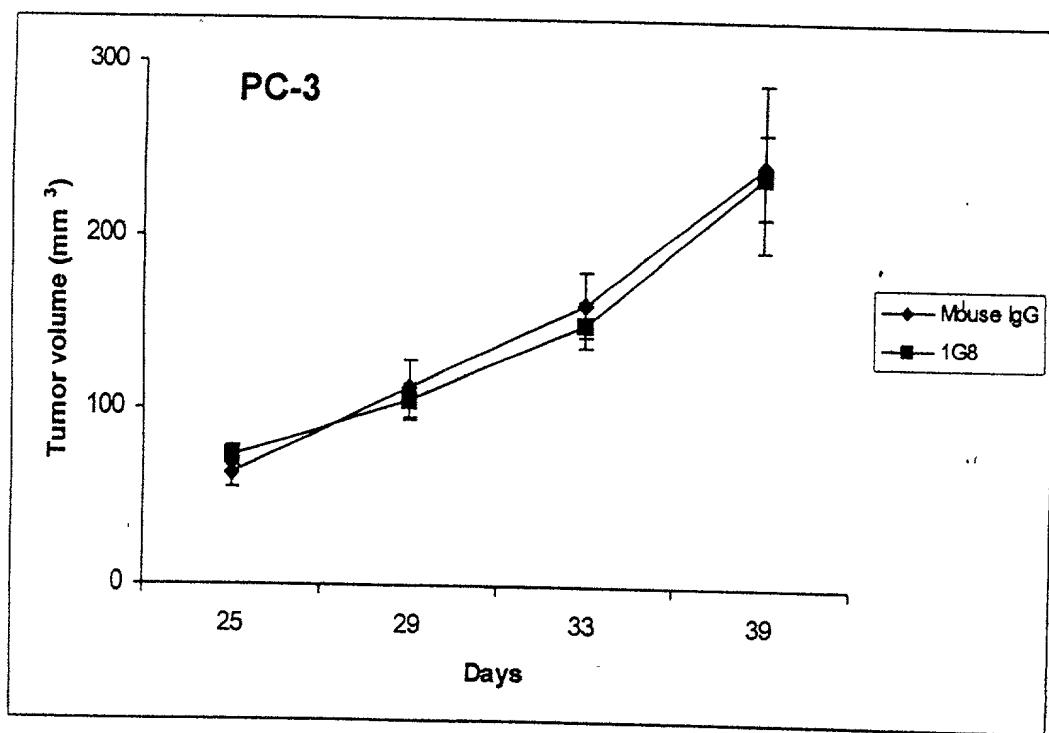
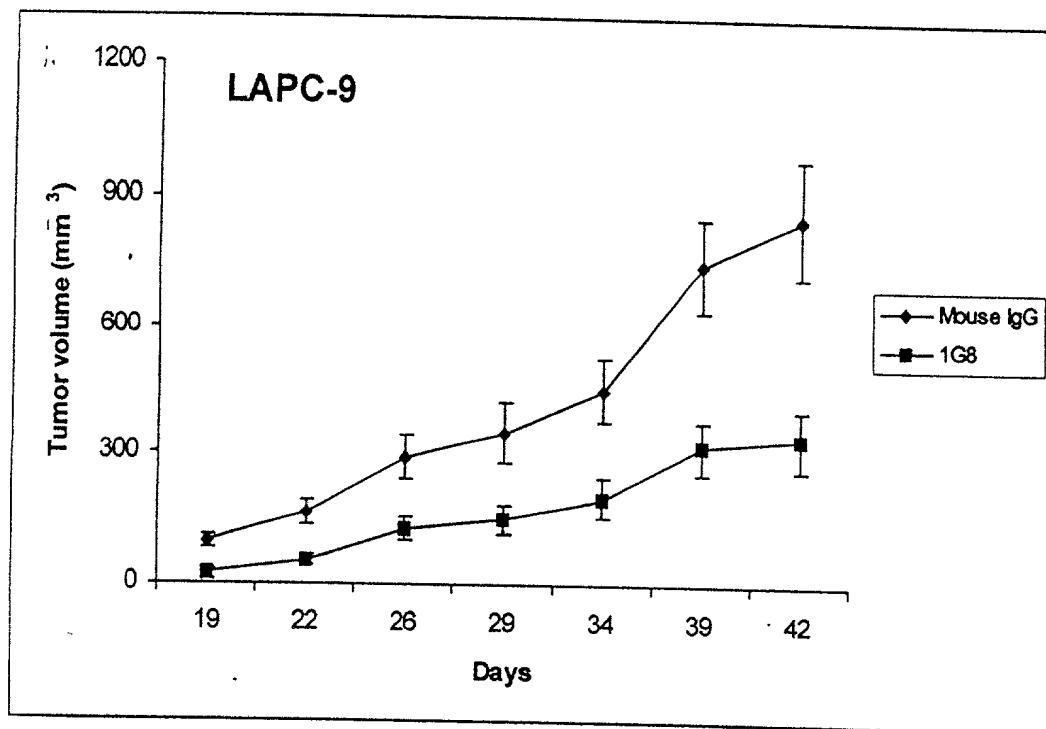
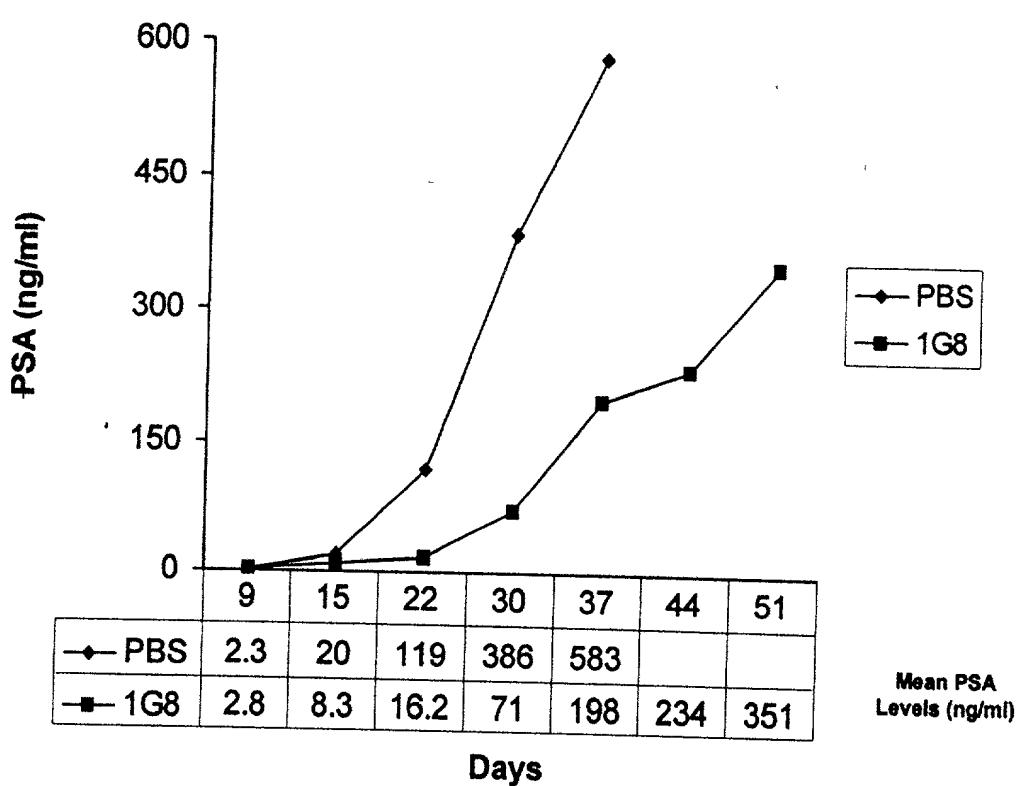


FIGURE 65

A)



B)

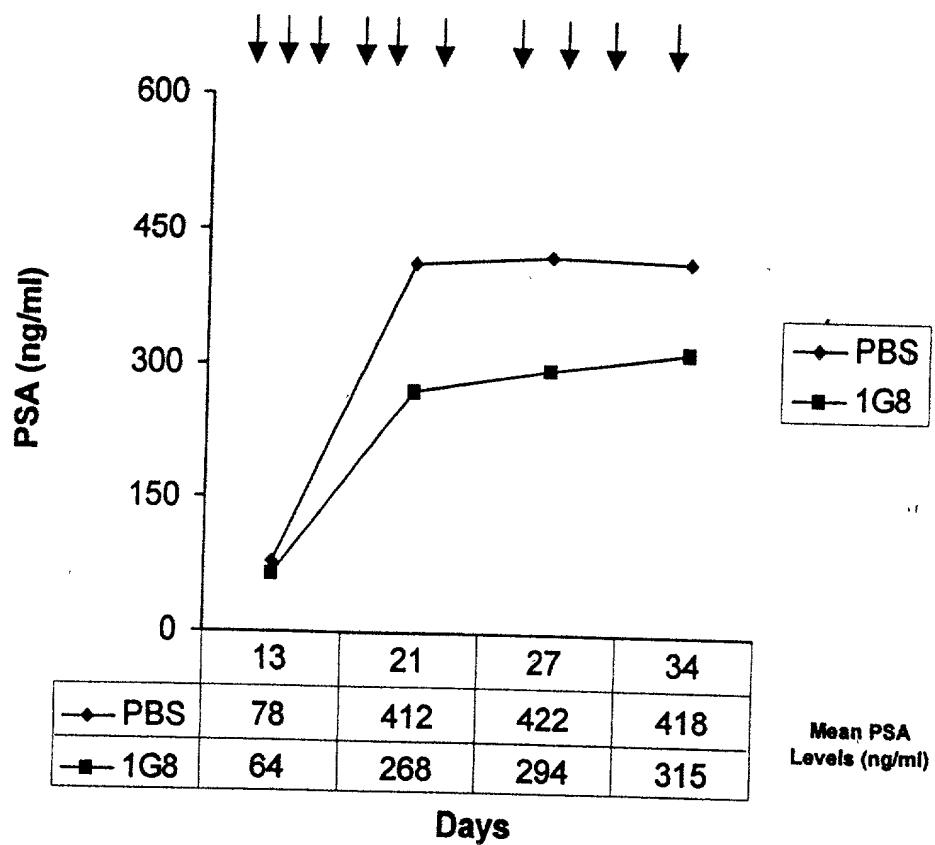
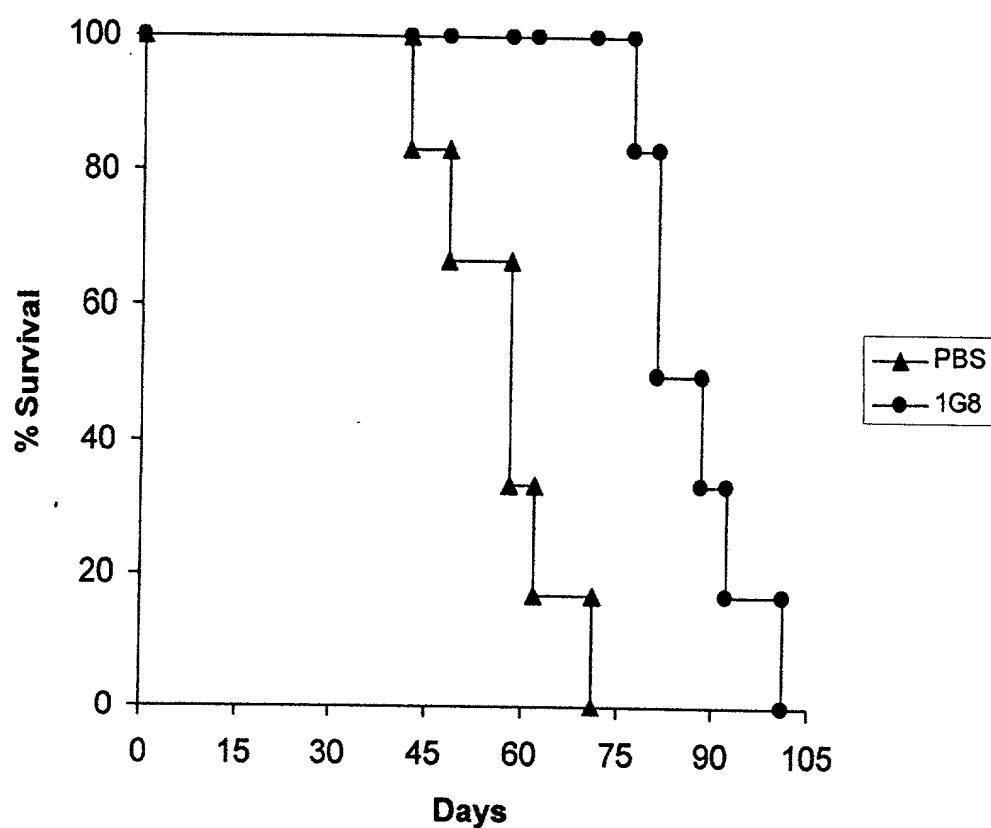


Figure 66

A)



B)

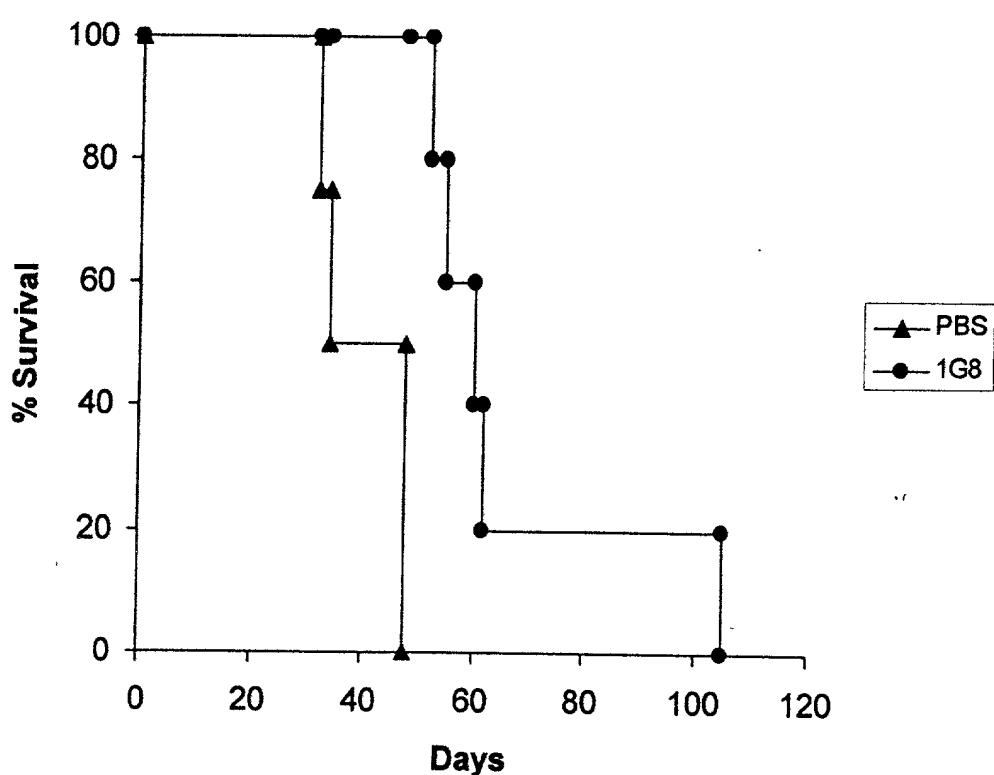


Figure 67

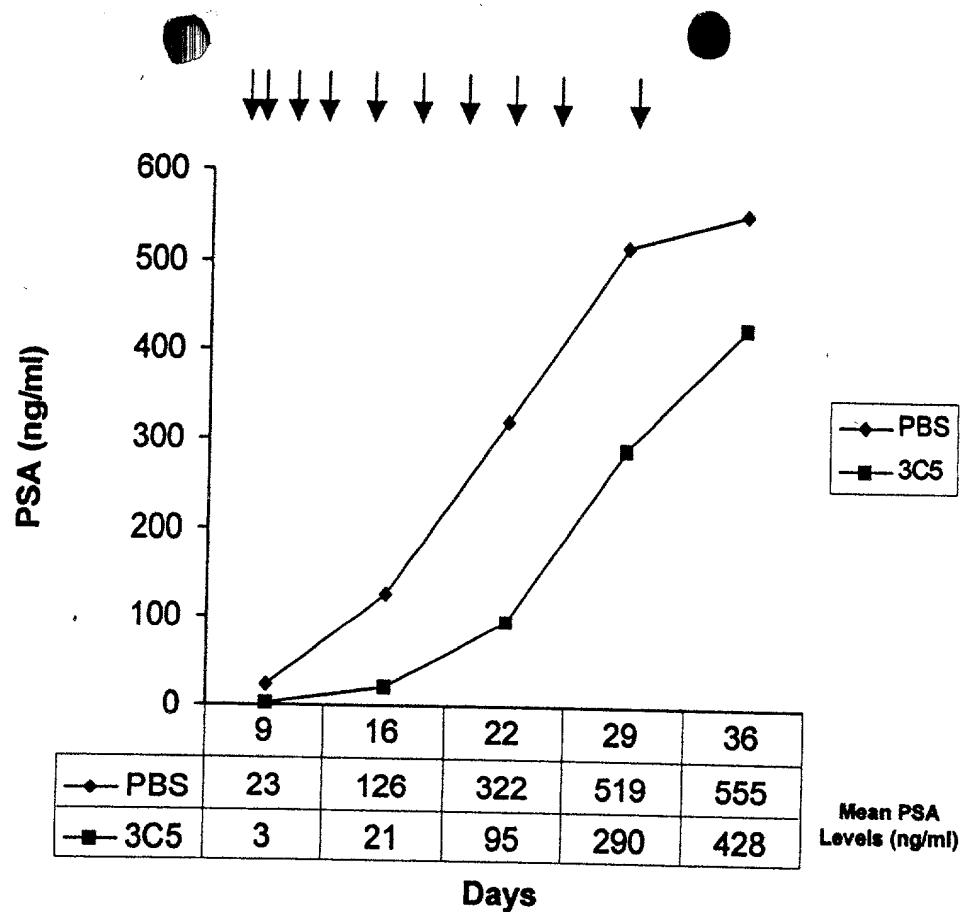
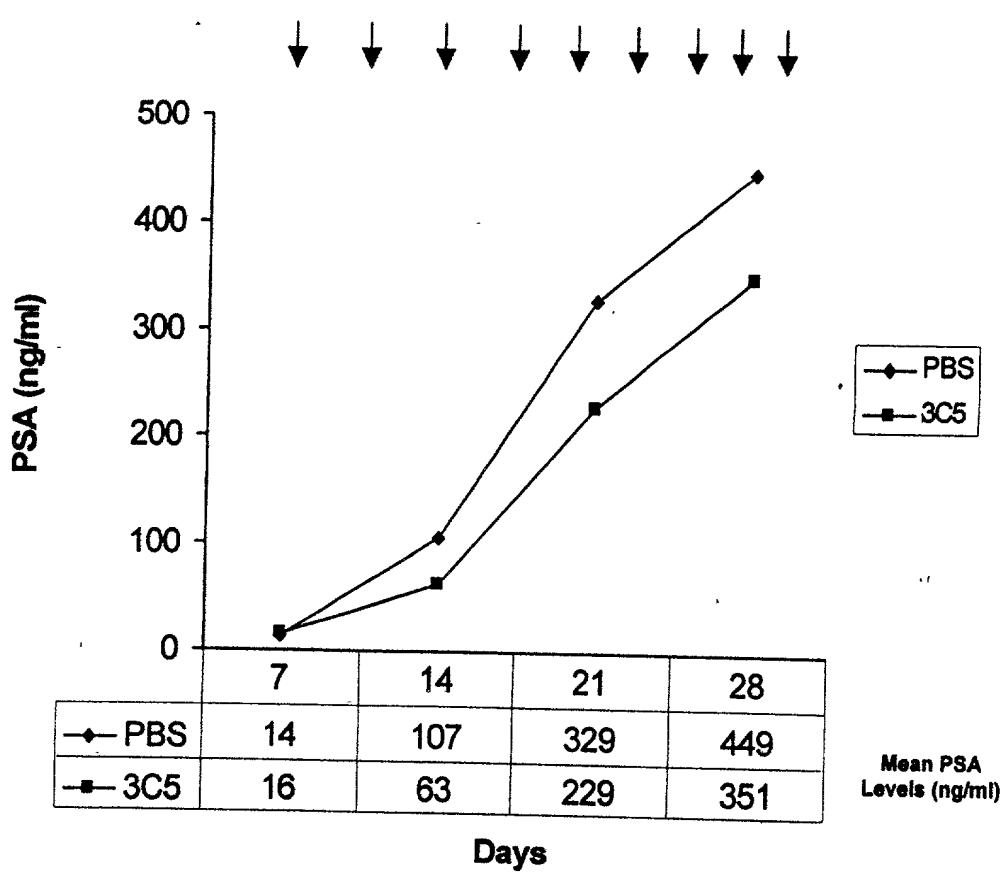
A)**B)**

Figure 68

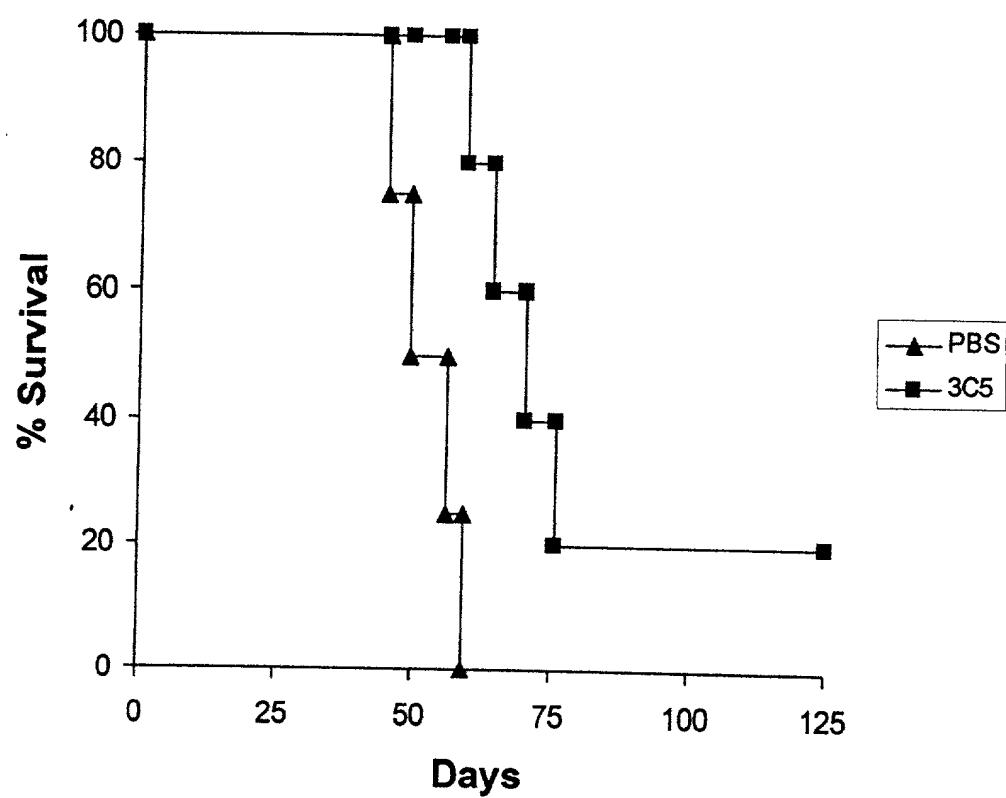
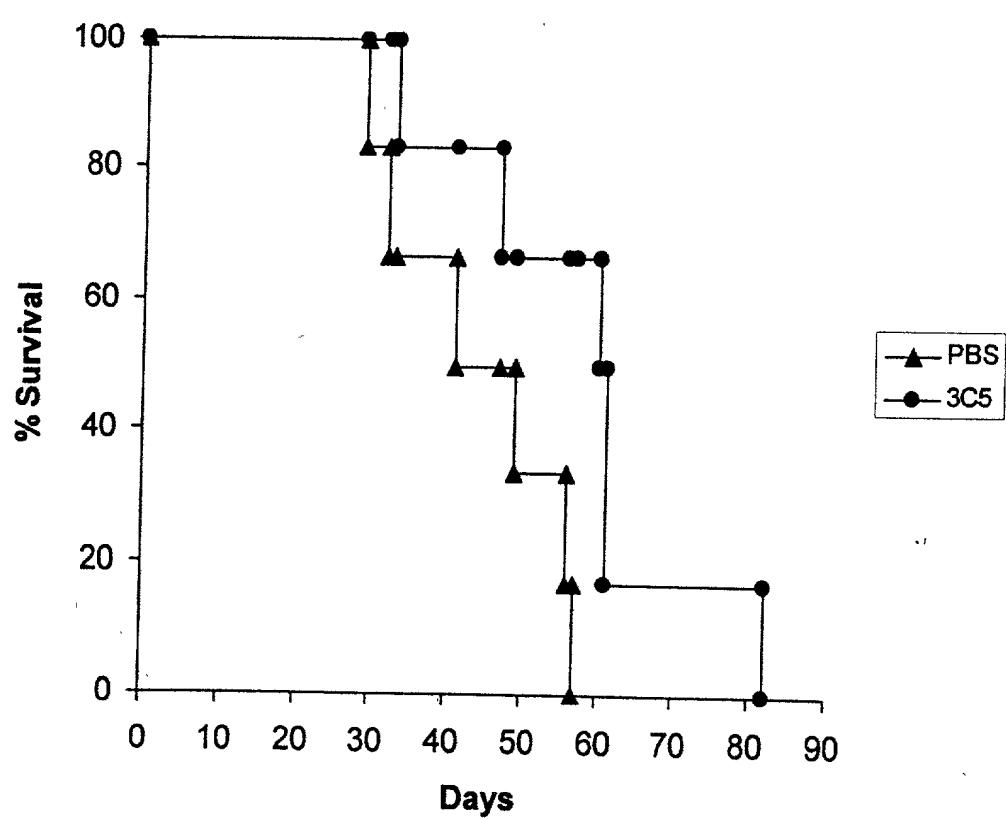
A)**B)**

Figure 69

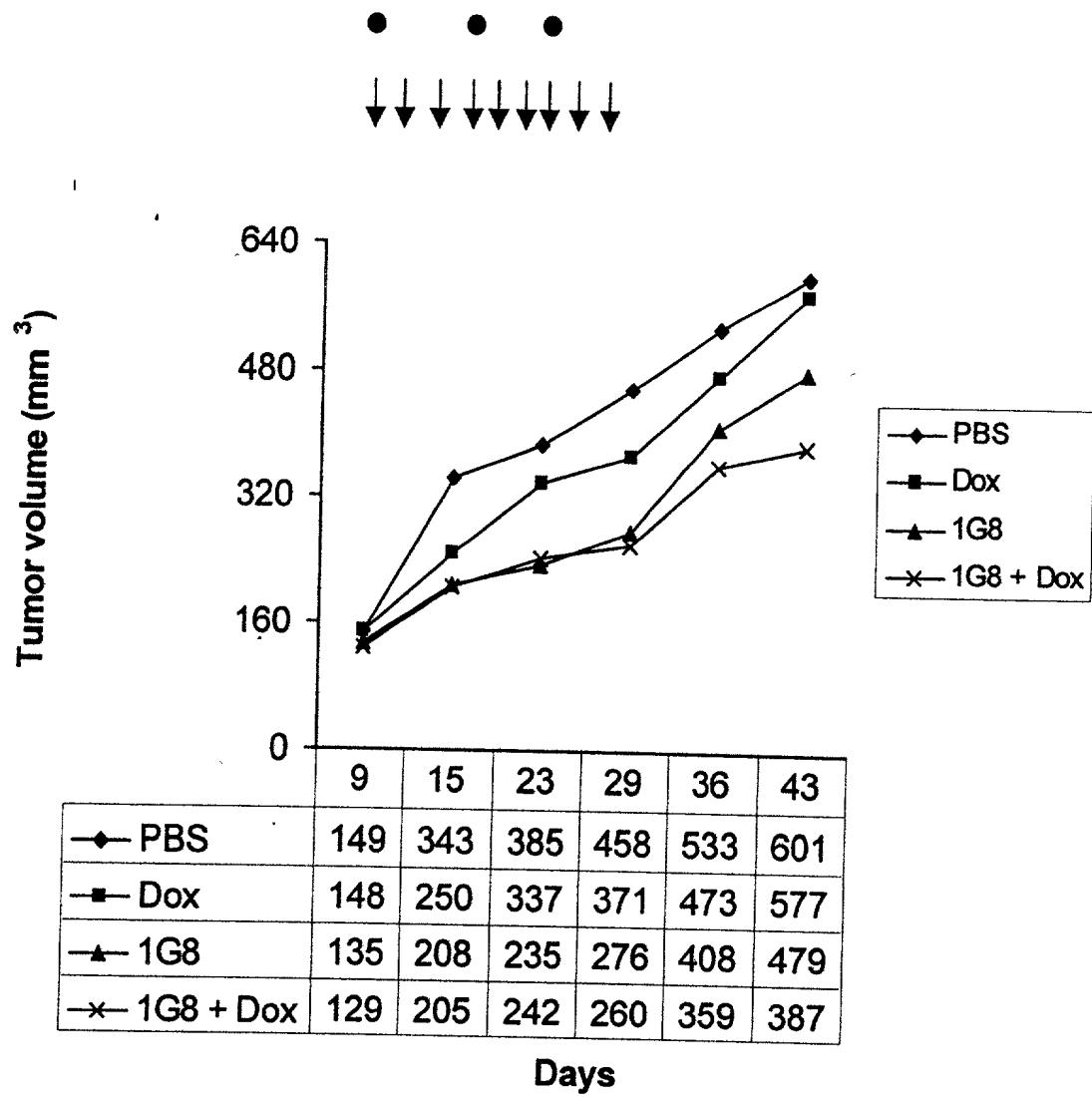
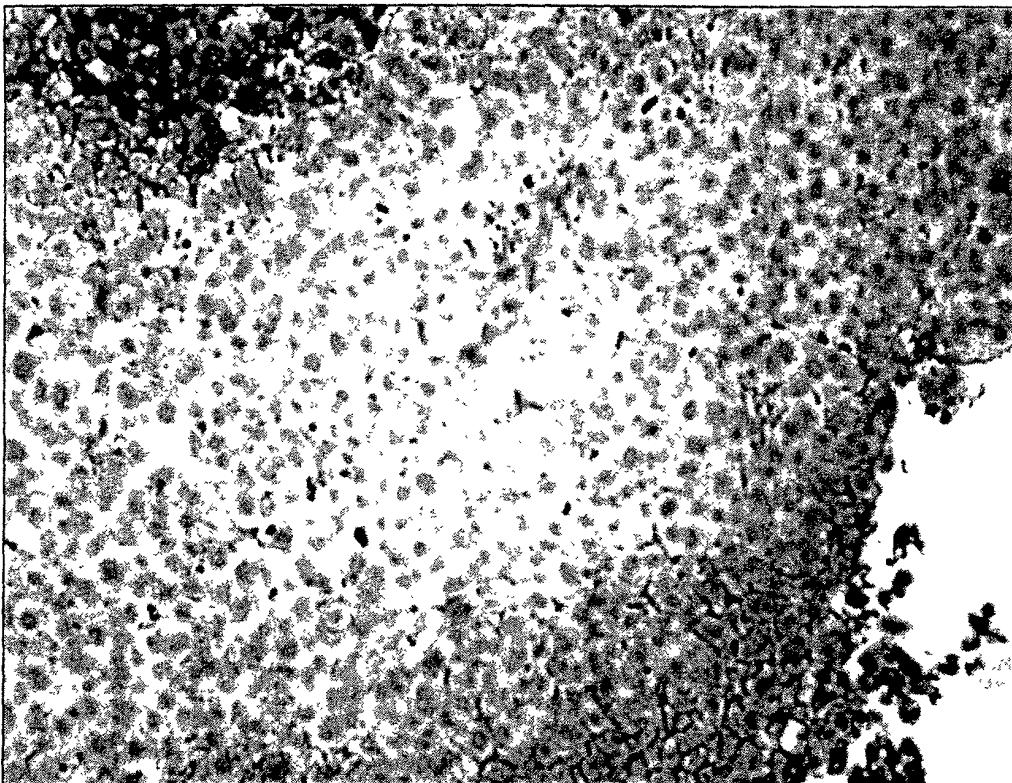


Figure 70

PSCA 3C5 MAb Localizes within LAPC9AD Xenograft Tissue

3C5 Treated



mlgG Treated

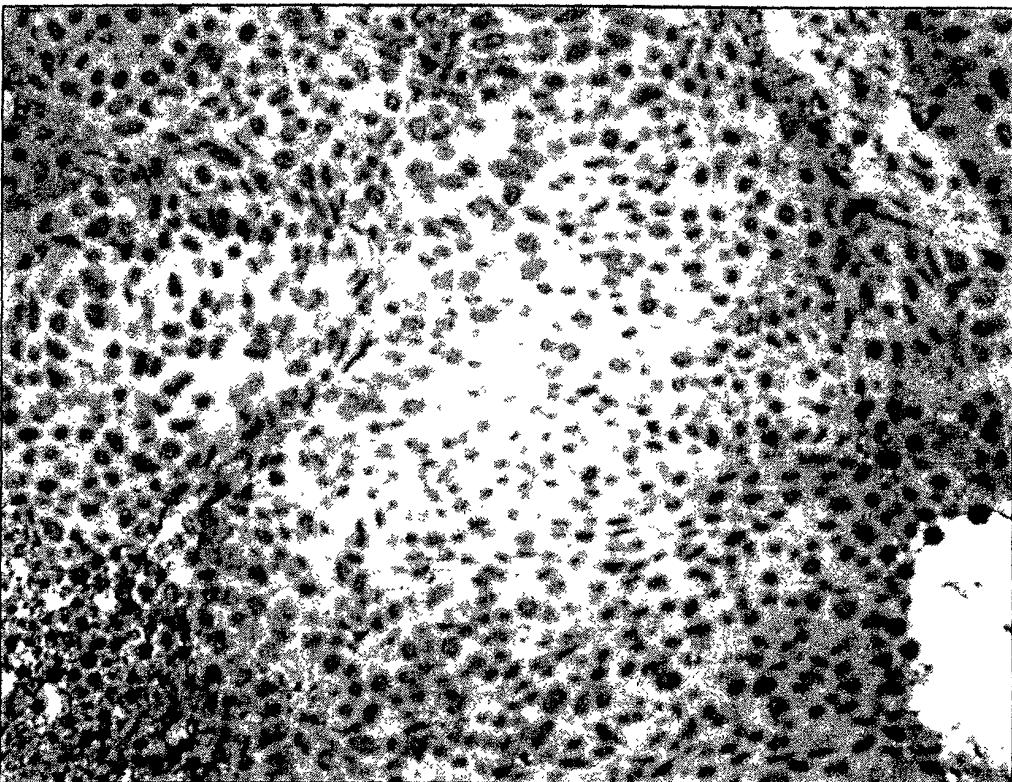


Figure 71

3C5 Anti-PSCA MAb is Localized to Established LAPC-9 Tumors

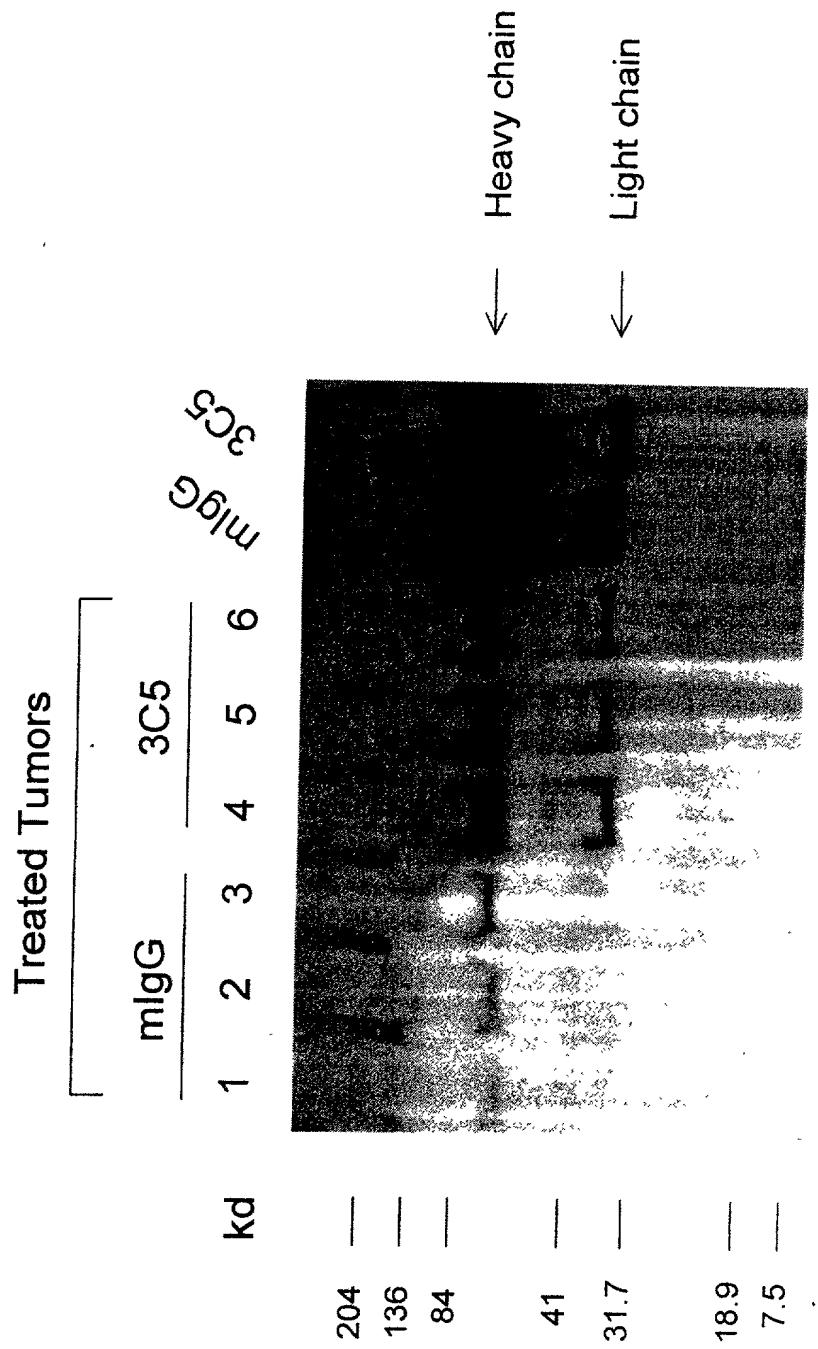
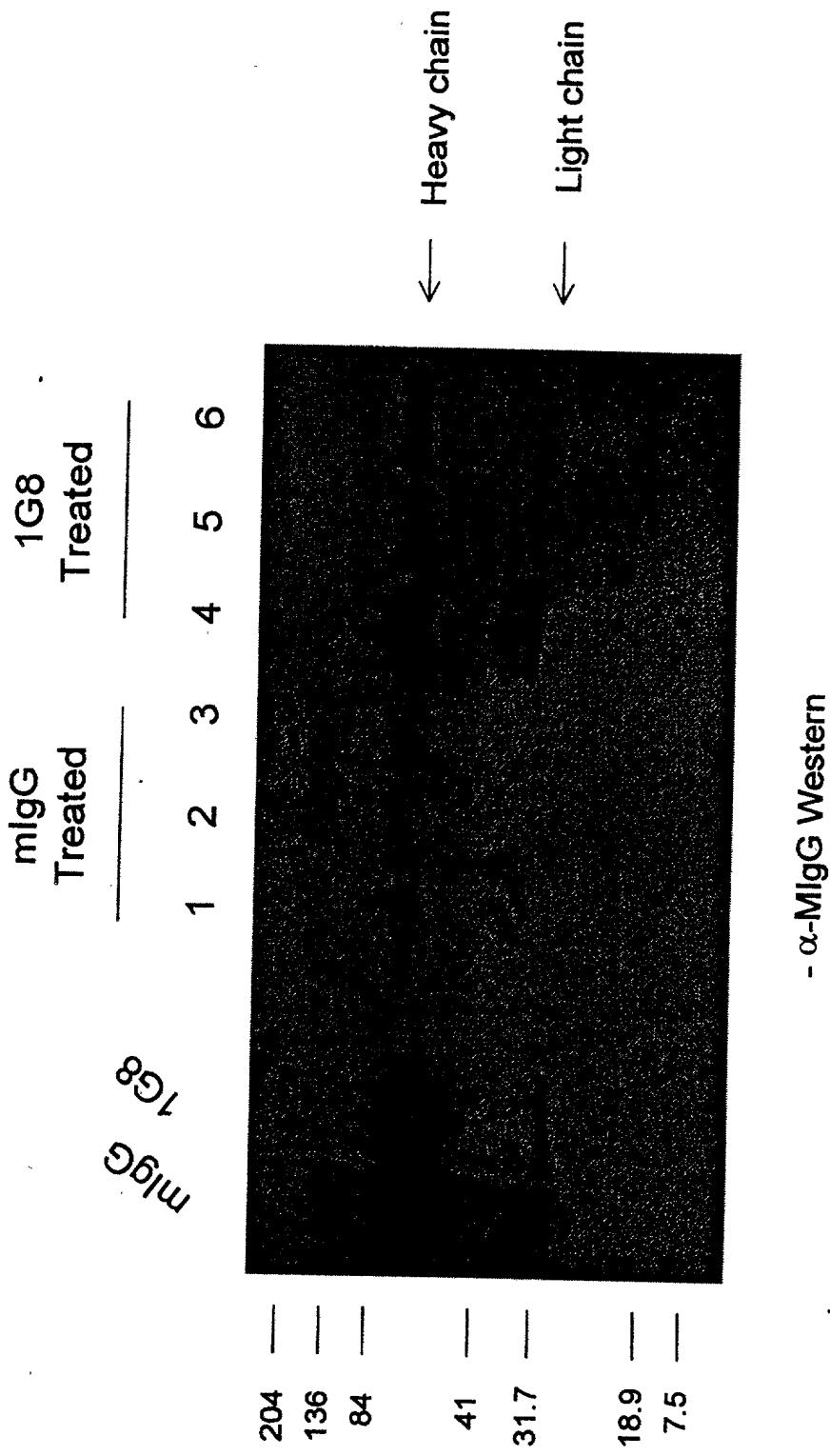


Figure 72

SPECIFIC TARGETING OF THE 1G8 ANTI-PSCA MAb TO ESTABLISHED LAPC-9 TUMORS



Method: Mice bearing established LAPC-9 tumors ($>100 \text{ mm}^3$) were injected with either mIgG or the anti-PSCA MAb 1G8. Tumors were harvested a week later and made into protein lysates for Western analysis.

Figure 73